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          WPI; 2003-247999/24.
P-PSDB; ABR63251.
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19-JUL-2001; 2001US-0306161P.
16-NOV-2001; 2001US-0331477P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel neural thread protein peptide, referred as cell death peptide, useful for treating prostatic hyperplasia, psoriasis, eczema, dermatos atherosclerosis, cosmetic modification to skin, throat, mouth, muscle.
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                                                                                                                                                                                                                                                                                                                              encoding sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP; 12 A; 20 C; 14 G; 11 T; 0 U; 0 Other;
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2001US-0331477P.
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                                                                                 CORP
                                                                                                                                                                                                                                                                                     protein; NTP; tumour; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                      DNA;
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100.0%; Pred. No.
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. 36;
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Novel neural thread protein peptide, referred as cell death peptide, useful for treating prostatic hyperplasia, psoriasis, eczema, dermatos atherosclerosis, cosmetic modification to skin, throat, mouth, muscle.
                                                                                                                                                                    dermatosis,
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Disclosure; Page 16; 77pp; English.

The present invention relates to a neural thread protein (NTP) peptide referred to as cell death peptide. Thought to be cytostatic, antibacterial, immunosuppressive and antiinflammatory. It is useful for treating a condition in a patient requiring removal or destruction of cells, for treating a condition such as benign or malignant tumor, inflammatory disease, autoimmune disease and infectious disease. The peptide useful for treatment is derived from the amino acid sequence for a pancreatic thread protein. The peptide is conjugated, linked or bound to a molecule chosen from antibody or its fragment, antibody-like binding molecule, where the molecule has a higher affinity for binding to a tumor or other target than binding to other cells. Treatment using NTP peptides can remove benign tumors with less risk and fewer of the undesirable side effects of surgery. The present sequence is an NTP encoding sequence

Sequence 57 BP; 9 P, 21 C; 15 G; 12 T; 0 U; 0 Other;

δ Query Match Best Local S Matches 57 Local Similarity Conservative 5.8%; 0, Score 57; Pred. No. ore 57; DB 1 ored. No. 36; Mismatches ۲, 0 Length 57; Indels 0 Gaps 0

닭 RESULT 3 AAK91064 standard; DNA; 66 990 CCTCCCGGGCTCAAGCGATTCTCCTGTCTCAGCCTCCCAAGCAGCTGGGATTACGGG 1046 -CCTCCCGGGCTCAAGCGATTCTCCTGTCTCAGCCTCCCAAGCAGCTGGGATTACGGG 57 ВP

05-NOV-2001 AAK91064; (first entry)

Human digestive system antigen genomic sequence SEQ ID NO:

Human; digestive system antigen; gene therapy; cancer; appendicitis; ulcerative colitis; infection; Hirschsprung's disease; chronic colitis; digestive system disorder; Meckel's diverticulum; ds.

WO200155314-A2

17-JAN-2001;

2001WO-US001324

04-FEB-2000; 24-FEB-2000; 02-MAR-2000; 16-MAR-2000; 17-MAR-2000; 18-APR-2000; 30-JUN-2000; 07-JUN-2000; 28-JUN-2000; 19-MAY-2000; 2000US-018464P.
2000US-0189874P.
2000US-0199874P.
2000US-0190076P.
2000US-0190076P.
2000US-020515P.
2000US-021647P.
2000US-021647P.
2000US-021647P.
2000US-021680P.
2000US-0217496P.
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2000US-0220963P.
2000US-0224518P. 2000US-0179065P

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Matches 60
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17-NO
                                                                                                                                                                                               The present invention provides the protein and coding sequences of number of human digestive system antigens. These can be used in the diagnosis, treatment and prevention of digestive system disorders, including cancer, Meckel's diverticulum, bacterial or parasitic infections, appendicitis, Hirschsprung's disease, chronic colitis ulcerative colitis. The present sequence is a genomic DNA fragment encoding a digestive system antigen of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                 Polynucleotides encoding diagnosing, treating, pro
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                                                                                                                                                                                                                                                                                                                                                                                                            digestive
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RESULT 4
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2000US-0179065P.
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                                                                                                                                                                                                  disorders include autoimmune diseases such as rheumatoid arthritis, clisorders include autoimmune diseases such as rheumatoid arthritis, cardiovascular disorders such as cardiac arrest, cerebrovascular disorders such as cardiac arrest, cerebrovascular disorders such as cardiac arrest, cerebrovascular clischaemia, nervous system disorders such as Alzheimer's disease, infections caused by bacteria, viruses and fungi, ocular disorders such as corneal infection, endocrine disorders such as premature labour and infertility, gastrointestinal disorders such as Crohn's disease, renal disorders such as glomerulonephritis and respiratory disorders such as asthma and pleurisy. The polypeptides can also be used to aid wound healing, to prevent skin aging due to sumburn, to manintain organs before transplantation, to regenerate tissues and in the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                            Query Match
Best Local Similarity
Matches 60; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sequences AAS31827-AAS32182 represent genomic DNA molecules, which enco the liver associated polypeptides of the invention. Liver associated polypeptides are useful in the polypeptides and their associated polypucteotides are useful in the diagnosis, treatment and prevention of various types of disorders in ediagnosis, treatment and prevention of various types of disorders in ediagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Isolated nucleic acid molecule encoding a human liver related protein is used in preventing, treating or ameliorating disorders of the liver particularly cancer of the liver.
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                                                                                                                                                                                                                                                                                                                                                                                                                                   humans, mice, rabbits, goats, horses, cats, dogs, chickens or sheep. A pathological condition can be determined by detecting the presence or absence of a mutation in a liver associated polynucleotide. The treate
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                                                                     TTTTTGTATTTCATTAGAGGCGGGGTTTCACCATATTTGTCAGGCTGGTCTCAAACTCC 1129
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29 - SEP - 2000;
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03 - CCT - 2000;
04 - CCT - 2000;
07 - CCT - 2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; liver antigen; liver disorder; hepatic disorder; infection; hepatitis; viral; parasitic; bacterial; fungal; inflammatory condition; cirrhosis; granulomatous hepatitis; toxin damage; drug damage; autoimmune disease; wilson's disease; primary biliary cirrhosis; neoplastic disorder; cancer; tumour, portal hypertension; gastrointestinal disorder; hepatitis; drug screening; gene therapy; chromosome mapping; forensic analysis; antibody preparation; hepatotropic; cytostatic; antiinflammatory; virucide; antibacterial; hepatotropic; cytostatic; antiinflammatory; virucide; antibacterial;
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                                                                                                                                                                                                                                                                                                                                                                                                        14-AUG-
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14-AUG-2000;
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26-JUL-2000,
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14-AUG-2000
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 2000US-022526P.
2000US-022526P.
2000US-022526P.
2000US-022570P.
2000US-022575P.
2000US-022575P.
2000US-022575P.
2000US-0229343P.
2000US-0229343P.
2000US-0229343P.
2000US-0229343P.
2000US-0229343P.
2000US-0229343P.
2000US-023945P.
2000US-023945P.
2000US-023636P.
2000US-023636P.
2000US-023636P.
2000US-023636P.
2000US-023636P.
2000US-0236370P.
2000US-0236370P.
2000US-0237039P.
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2000US-0237039P.
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2000US-0220964P.
2000US-0224518P.
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RESULT 6
ADJ15367
ID ADJ1
XX
AC ADJ1
XX
                                                                                                                                                                                                                                                                                                                                                               crecombinant vectors and host cells comprising human liver antigen
CC plynucleotides, antibodies against human liver antigens, and the use of
CC liver antigen polynucleotides and polypeptides in diagnosing, treating,
CC prognosing or preventing various disorders of the liver. Such conditions
CC include viral infections (e.g., cytomegalovirus, Epstein-Barr virus,
CC infections (e.g., Clonorchis sinensis, Echinococcus granulosus and
CC Entamoeba histolytica), and also bacterial and fungal infections. Other
CC cirrhosis and granulomatous hepatitis), damage caused by drugs or toxins,
CC autoimmune diseases (e.g., Wilson's disease, primary biliary cirrhosis),
CC accinnoma), portal hypertension, or gastrointestinal disorders (e.g.,
CC peptic ulcers, gastritis and peritoneal diseases). Liver antigen
CC polypeptides and polynucleotides may also be used in screening for
CC compounds which modulate liver antigen expression or activity. The
CC polynucleotides may be used for gene therapy, chromosome mapping,
CC in the identification of individuals and in forensic analysis, and the
CC equences Newsons bused as molecular weight markers or to prepare
CC sequences Newsons bused as molecular weight markers or to prepare
CC sequences Newsons and the sequence data for this patent did not form part of
CC the printed specification, but was obtained in electronic format directly
CC from the USPTO at sequence captors obtained in electronic format directly
CC from the USPTO at sequence sequence of the printed format directly
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20-OCT-2000;
20-OCT-2000;
01-NOV-2000;
17-NOV-2000;
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08-DEC-2000;
08-DEC-2000;
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                   ADJ15367;
                                                  ADJ15367 standard; DNA; 66
                                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to 145 novel human liver antigens (ABP40831-ABP40975) and to cDNAs encoding them (ABN90036-ABN90180), and also encompasses polypeptides 90% identical and polynucleotides 95% identical to the sequences of the invention. The invention additionally relates to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; SEQ ID NO 575; 181pp; English.
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RUBEN S
BARASH
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2000US-0241785P.
2000US-0241809P.
2000US-0244617P.
2000US-0249299P.
2000US-0251868P.
2000US-0251868P.
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Pred. No. 44;
0; Mismatches
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hepatic cancer, also
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   18-AUG-2000;
22-AUG-2000;
22-AUG-2000;
23-AUG-2000;
30-AUG-2000;
01-SEP-2000;
01-SEP-2000;
01-SEP-2000;
01-SEP-2000;
05-SEP-2000;
06-SEP-2000;
06-SEP-2000;
08-SEP-2000;
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30-JUN-2000;
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В 5

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liver; virucide; fungicide; antibacterial; antiparasitic; hepatotropic; antiinflammatory; cytostatic; litholytic; antirheumatic; antiarthritic; neuroprotective; antidabetic; anticoagulant; thrombolytic; antiarteriosclerotic; cardiant; haemostatic; antiarrhythmic; ophthalmological; antiarteriosclerotic; vasotropic; osteopathic; nootropic; antiparkinsonian; anticonvulsant; neuroleptic; vasotropic; orotropic; antiparkinsonian; anticonvulsant; neuroleptic; vasotropic; cytostatic; gynaecological; viral; fungal; bacterial; parasitic infection; citrhosis; Wilson's disease; gastrointestinal disorder; pancreatic; gallbladder; immune; blood; hyperproliferative; cardiovascular; respiratory; musculoskeletal system; neurological; endocrine; reproductive system; developmental; inherited; hyperproliferative; cardiovascular; respiratory; musculoskeletal system;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human liver-related genomic
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2000US-022964P 2000US-0224518P 2000US-0225214P 2000US-0225214P 2000US-0225266P 2000US-0225266P 2000US-0225267P 2000US-0225247P 2000US-0225759P 2000US-0225759P 2000US-0225759P 2000US-022679P 2000US-0229349P 2000US-0229349P 2000US-0229349P 2000US-0229349P 2000US-0239343P 2000US-0239343P 2000US-0239343P 2000US-0239343P 2000US-023943P 2000US-023943P 2000US-023943P 2000US-023943P 2000US-023943P 2000US-023943P 2000US-023943P 2000US-0216880P. 2000US-0217487P. 2000US-0217496P. 2000US-0218290P. 2000US-0214886P. 2000US-0215135P. 2000US-0216647P. 2002US-00073961 2000US-0205515P. 2000US-0209467P. 2000US-0198123P 2000US-0190076P 2000US-0189874P 2000US-0184664P 2000US-0179065P

08-SEP-2000; 08-SEP-2000; 08-SEP-2000; 12-SEP-2000; 14-SEP-2000; 14-SEP-2000; 14-SEP-2000;

14-SEP-2000; 14-SEP-2000; 14-SEP-2000; 14-SEP-2000;

2000US-0234997P. 2000US-0234998P.

2000US-0231414P.
2000US-0232080P.
2000US-0232080P.
2000US-0231968P.
2000US-0232399P.
2000US-0232399P.
2000US-0232400P.
2000US-0232400P.
2000US-0233400P.
2000US-0233403P.
2000US-0233403P.
2000US-0233063P.
2000US-0233064P.
2000US-0233064P.
2000US-02334274P.

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RESULT 7
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AC ADI2
XC ADI2
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                                                                                                                                                                                                                                                                                                                                                                                                                cc osteopathic, nootropic, antiparkinsonian, anticonvulsant, neuroleptic, costopathic, nootropic, and gynaecological activities. The polypoptides cc and polynucleotides of the invention may be useful for diagnosis, cc detection, treatment and/or prevention of disorders of the liver such as contract the state of the liver such as contract the specification per se but was obtained electronically from the USPTO constitute.
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Best Local S
Matches 60
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17.NOV-2000;
01.DEC-2000;
01.DEC-2000;
05.DEC-2000;
05.DEC-2000;
06.DEC-2000;
06.DEC-2000;
08.DEC-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to a novel isolated, liver related polypeptide. The polypeptide of the invention demonstrates virucide, fungicide, antibacterial, antiparasitic, hepatotropic, antiinflammatory, cytostatic, litholytic, antirheumatic, antiarthritic, neuroprotective, antidiabetic, anticoagulant, thrombolytic, antiarteriosclerotic, cardiant, haemostatic, antiarrhythmic, ophthalmological, antiarteriosclerotic, vasotropic, antiarrhythmic, ophthalmological, antiarteriosclerotic, vasotropic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New liver related polypeptide, useful for diagnosis, treatment and/or prevention of liver, gastrointestinal, pancreatic, immune, blood related, endocrine, reproductive, hyperproliferative or reproductive disorders.
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Oligonucleotide sequence enquiry #60.
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| 60; Conserv
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                                                                                                        standard; DNA; 60
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nilarity 90.9%;
Conservative (
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2000US-0251879P.
2000US-0251868P.
2000US-0251869P.
2000US-0251969P.
2000US-0251909P.
2000US-0254097P.
2001US-0259678P.
2001US-00764887.
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2000US-024930P.
2000US-0250160P.
2000US-0250391P.
2000US-0251939P.
2000US-0251988P.
2000US-0256719P.
                                       (first entry)
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14-SEP-2000;
21-SEP-2000;
21-SEP-2000;
25-SEP-2000;
26-SEP-2000;
27-SEP-2000;
29-SEP-2000;
29-SEP-2000;
29-SEP-2000;
29-SEP-2000;
29-SEP-2000;
29-SEP-2000;
29-SEP-2000;
20-OCT-2000;
21-OCT-2000;
21-OCT-2000;
20-OCT-2000;
20-OC

2000US-0235484P.
2000US-0235834P.
2000US-0235837P.
2000US-0235359P.
2000US-0235369P.
2000US-023569P.
2000US-023569P.
2000US-023690P.
2000US-023703P.
2000US-023703P.
2000US-023703P.
2000US-023703P.
2000US-0240960P.
2000US-0241785P.
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2000US-0241787P.
2000US-0241787P.
2000US-0246778P.
2000US-0246477P.
2000US-0246478P.
2000US-0246478P.
2000US-0246528P.

17-NOV-2000; 17-NOV-2000;

2000US-024660P 2000US-0246611P 2000US-0246611P 2000US-024920P 2000US-024920P 2000US-024920P 2000US-0249210P 2000US-0249211P 2000US-0249211P 2000US-0249213P 2000US-0249213P 2000US-0249213P 2000US-0249213P 2000US-0249214P 2000US-0249214P 2000US-0249214P 2000US-0249217P 2000US-0249214P 2000US-0249217P 2000US-0249249P 2000US-0249249P 2000US-0249249P 2000US-0249249P 2000US-0249265P 2000US-0249265P

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Best Local S
Matches 57
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         comprises identifying non-protein-encoding nucleotide sequences within an mRNA transcript or a DNA sequence encoding same in the nucleome. The methods are useful for identifying an eRNA or DNA for modifying a genetic network in cell to alter the cells phenotype. The present sequence represents human oligonucleotide sequence enquiry.
Scholl T,
                                             07-JUN-2002;
09-AUG-2002;
                                                                                                                               WO2003104474-A2
                                                                                                                                                                           ds; cancer; human; tumour suppressor;
breast cancer susceptibility gene 1; BRCA1; repetitive Alu;
ovarian cancer; recombination; mutant.
                                                                                                                                                                                                                             Mutant
                                                                                                                                                                                                                                                                                                  ADI12552 standard; DNA; 66
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention relates to identifying an eRNA or a DNA sequence comprising an eRNA-encoding sequence in the nucleome of a eukaryotic \sigma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Identifying an eRNA or a DNA sequence comprising an eRNA-encoding sequence in the nucleome of a eukaryotic cell, comprising identifying process-protein-encoding nucleotide sequences within an mRNA transcript or a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19-SEP-2001; 2001US-0324127P
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                                                                                09-JUN-2003; 2003WO-US018098
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                       (MYRI-) MYRIAD GENETICS INC.
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                                                                                                                                                                                                                                                                                                                                                                                  CTCTGTCACCCAGGCTGGAGTGCAGTGGCGCAATCTTGGCTCACTGCAACCTCTGCCTCC
Hendrickson BC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP; 8 A; 22 C; 16 G; 14 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQ ID NO 63; 137pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                             2002US-0387132P.
2002US-0402430P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2002WO-AU001286
                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                            genomic
                                                                                                                                                                                                                                                                                                                                                                                                                    5.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Stanley S;
                                                                                                                                                                                                                                                                                                  BP.
                                                                                                                                                                                                                            DNA resulting
                                                                                                                                                                                                                                                                                                                                                                                                          <u>,</u>
 Ward B,
                                                                                                                                                                                                                                                                                                                                                                                                                   Score 55.2;
Pred. No. 47
                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
 Pruss D;
                                                                                                                                                                                                                             from
                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1; Length 60;
                                                                                                                                                                                                                                                                                                                                                                                                         ω
                                                                                                                                                                                                                            deletion 5 SeqID
                                                                                                                                                                                                                                                                                                                                                                                                          Indels
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                                                                                                                                                                                                                             <u>ω</u>
                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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DNA
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RESULT 9
ADI12619/c
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Best Local S
Matches 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              present invention describes newly discovered deletion mutations that are believed to be deleterious and cause significant alterations in the structure or biochemical function of BRCA1. Accordingly, it provides methods for detecting such mutants, as well as identifying and screening for cytostatic compounds useful for treating or preventing cancers associated with a BRCA1 genetic variant. This polynucleotide is a mutant human BRCA1 genomic DNA fragment that arrises as a result of a recombination event (deletion 5), which causes the omission of exons 15 and 16, given in an exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This invention relates to a novel method for predicting a predispositi to cancer in a patient by detecting large deletions in the human tumou suppressor gene identified as the breast cancer susceptibility gene 1 (BRCA1). Specifically, it refers to deletions that result from the unequal crossover between a pair of repetitive Alu sequences in the gene, such that the recombined nucleotide sequence containing the deletion indicates a predisposition to breast and ovarian cancer. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Predicting a predisposition to cancer in a patient comprising detecting deletion in the BRCA1 gene that results from the unequal crossover between a pair of repetitive sequences in the BRCA1 gene.
                                                                                                                                                                                                                                          07-JUN-2002;
09-AUG-2002;
                                                                                                                                                                                                                                                                                                                                    09-JUN-2003; 2003WO-US018098
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ds; cancer; human; tumour suppressor;
breast cancer susceptibility gene 1;
ovarian cancer; recombination.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human BRCA1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO2003104474-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
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                                                                                                                                                                                (MYRI-) MYRIAD GENETICS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CCGCCTCCCGGGTTCAAGCAATTCTCCTGCCTCAGCCTCCTGAGTAGCTGGGATTACAGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 66 BP; 15 A; 16 C; 24 G; 11 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5.5%;
nilarity 90.6%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA upstream from the deletion
                                                                                                                                                                                                                                          2002US-0387132P
2002US-0402430P
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Pred. No. 55;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BRCA1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  5 recombination event.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  repetitive Alu;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       9
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Predicting a predisposition to

cancer in a patient comprising detecting

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WPI; 2004-062369/06

Hendrickson

BC,

Ward

B

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RESULT 10
AAK86585
ID AAK86
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Best Local S
Matches 58
 31-JAN-2000
04-FEB-2000
24-FEB-2000
02-MAR-2000
16-MAR-2000
17-MAR-2000
17-MAR-2000
19-MAY-2000
07-JUN-2000
07-JUN-2000
07-JUL-2000
07-JUL-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This invention relates to a novel method for predicting a predisposition to cancer in a patient by detecting large deletions in the human tumour suppressor gene identified as the breast cancer susceptibility gene 1 (BRCA1). Specifically, it refers to deletions that result from the unequal crossover between a pair of repetitive Alu sequences in the BRCA1 gene, such that the recombined nucleotide sequence containing the deletion indicates a predisposition to breast and ovarian cancer. The present invention describes newly discovered deletion mutations that are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     believed to be deleterious and cause significant alterations in the structure or biochemical function of BRCA1. Accordingly, it provides methods for detecting such mutants, as well as identifying and screening for cytostatic compounds useful for treating or preventing cancers associated with a BRCA1 genetic variant. This polynucleotide is a human BRCA1 DNA fragment representing the region downstream of the deletion 5 recombination event that causes the omission of exons 15 and 16, given in an exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Fig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               deletion in the between a pair of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                       07-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAK86585 standard; DNA; 63 BP
                                                                                                                                                                                                                                        17-JAN-2001; 2001WO-US001354.
                                                                                                                                                                                                                                                                     09-AUG-2001.
                                                                                                                                                                                                                                                                                                   WO200157182-A2
                                                                                                                                                                                                                                                                                                                                                             cytostatic; gene therapy; vaccine; metastasis;
                                                                                                                                                                                                                                                                                                                                                                              Human;
                                                                                                                                                                                                                                                                                                                                                                                                         Human immune/haematopoietic antigen genomic sequence SEQ ID NO:41397
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    687
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                58;
                                                                                                                                                                                                                                                                                                                                                                  immune; haematopoietic; immune/haematopoietic antigen; cancer
atic; gene therapy; vaccine; metastasis; ds.
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; 2000US-0179065P,
2000US-0184664P,
2000US-0186350P,
2000US-0186350P,
2000US-0190076P,
2000US-0190076P,
2000US-0295515P,
2000US-0205515P,
2000US-0214886P,
2000US-0215135P,
2000US-0216847P,
2000US-0217487P,
2000US-0217486P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 54.4; DI
Pred. No. 55;
0; Mismatches
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       14-AUG-2000)
12-AUG-2000)
12-AUG-2000)
13-AUG-2000)
10-SEP-2000)
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2000US-0229874P.
2000US-0229344P.
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2000US-0229349P.
2000US-0230438P.
2000US-0231242P.
2000US-0231243P.
2000US-0231244P.
2000US-0231244P.
2000US-0231244P.
2000US-0231244P.
2000US-0231244P.
2000US-0231249P.
2000US-0231414P.
2000US-023298P.
2000US-023299P.
2000US-0232399P.
2000US-02333064P.
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2000US-0220964P.
2000US-0220964P.
2000US-0224518P.
2000US-0225211P.
2000US-0225214P.
2000US-0225266P.
2000US-0225268P.
2000US-0225270P.
2000US-0225277P.
                    2000US-0232401P
2000US-0233064P
2000US-0233065P
2000US-0234274P
2000US-0234299P
2000US-0235484P
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2000US-0235834P
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2000US-023632P
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2000US-0241809P
2000US-0241809P
2000US-0241809P
2000US-0241809P
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2000US-0225759P.
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2000US-0246474P 2000US-0246475P

Indels

<u>,,</u>

Gaps

0

61

63;

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amino acid sequences given in AAM82170 to AAM91921. (I) have cytostatic activity, and can be used in gene therapy and vaccine production. (I) proteins and polynucleotides may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate (I) expression. For example, they may be used to treat disorders associated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of (I) by expressing inactive proteins or to supplement the patients own production of (I). Additionally, (I) polynucleotides may be used to produce the secreted (I), by inserting the nucleic acids into a host cell and culturing the cell to express the protein. (I) proteins and polynucleotides may be used to prevent, diagnose and treat immune/haematopoietic-related diseases, especially
                                                                                                                                                                                                                                                                             AAK54951 to AAK64702 encode the human immune/haematopoietic antigen amino acid sequences given in AAM82170 to AAM91921. (I) have cytosta
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2000US-0246528P

2000US-0246528P

2000US-0246528P

2000US-0246532P

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2000US-0246611P

2000US-0249207P

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2000US-0249219P

2000US-0249219P

2000US-0250160P

2000US-0251968P

2000US-0251868P

2000US-0251868P
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2000US-0251990P.
2000US-0254097P.
2001US-0259678P.
                                                                                                                                                                                                                                                                                                                                                     41397; 3071pp + Sequence Listing; English
                                                                                                                                                                                                                                                                                                                                                                                                  human immune/hematopoietic diagnosing and/or treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MS
                                                                                                                                                                                                                                                                                                                                                                                                    : antigen
                                                                                                                                                                                                                                                                                                                                                                                                  polypeptides, and metastasis.
            RESULT: 11
AAK85681/c
ID AAK856
XX AAK856
XX AAK856
XX Human
XX Human;
XW Human;
XW Cytost
XX O9-AUG
XX 17-JAN
PR 11-AUG
PR 11
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                                                                                                          04-FEB-2000
24-FEB-2000
12-MAR-2000
16-MAR-2000
19-MAR-2000
19-MAY-2000
28-JUN-2000
29-JUN-2000
30-JUN-2000
07-JUL-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
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ilarity 91.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       therapy; vaccine; metastasis;
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antigen;

cancer;

SEQ ID NO: 40493.

Disclosure;

SEQ ID

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Nucleic

acids encoding for preventing,

2001-483426/52

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Barash

Ruben

HUMAN

GENOME SC,

SCI

08-NOV-2000
01-NOV-2000
017-NOV-2000

22-AUG-2000; 22-AUG-2000; 23-AUG-2000; 30-AUG-2000; 01-SEP-2000; 01-SEP-2000; 01-SEP-2000; 05-SEP-2000; 06-SEP-2000; 06-SEP-2000; 08-SEP-2000; 08-SEP-2000;

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2000US-0231242P.
2000US-0231413P.
2000US-0231968P.
2000US-0232399P.
2000US-0232399P.
2000US-023239P.

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                                                                                                                                                                 AAKS4951 to AAK64702 encode the human immune/haematopoietic antigen (I) CC amino acid sequences given in AAM82170 to AAM91921. (I) have cytostatic CC activity, and can be used in gene therapy and vaccine production. (I) CC proteins and polynucleotides may be used in the prevention, diagnosis and CC treatment of diseases associated with inappropriate (I) expression. For CC example, they may be used to treat disorders associated with decreased CC expression by rectifying mutations or deletions in a patient's genome CC supplement the activity of (I) by expressing inactive proteins or to CC supplement the patients own production of (I). Additionally, (I) CC supplement the patients own production of (I). Additionally, (I) CC diagnose and treat immune/haematopoietic acids into a host cell and culturing the cell to express the CC protein. (I) proteins and polynucleotides may be used to prevent, CC diagnose and treat immune/haematopoietic-derived cells. AAK64703 CC cancers and cancer metastases of haematopoietic antigen genomic CC sequences from the present invention. AAK54942 to AAK54950 and AAM82169 crepresent sequences used in the exemplification of the present invention.
                                                                                      Matches
                                                                                                  Query Match
Best Local
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17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
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05-DEC-2000;
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17-NOV-2000;
17-NOV-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; SEQ ID NO 40493; 3071pp + Sequence Listing; English
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                                                                                                                                            Sequence
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   232 CA 233
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                              62
                                                                                    l Similarity
57; Conserv
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                                                    TTTTTTAGTAGAGATGGAGTTTCTCCATGTTGGTCAGGCTGGTCTCGAACTCCCGACCT
                                                                                                                                              63
                            Barash
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2000US-0251030P.

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2000US-0256719P.

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2000US-0251869P.

2000US-0251989P.

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2000US-0249218P.
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                                                                                      Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 human immune/hematopoietic antigen polypeptides, diagnosing and/or treating cancers and metastasis.
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                                                                                    Score 54; DB Pred. No. 56; 0; Mismatches
                                                                                                                                              14 G; 11 T;
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                                                                                                                                               0 Other;
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                                                                                                                   Length 63;
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RESULT 12
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ID AAK83961;
XX
AX83961;
XX
AX83961;
XX
AX83961;
XX

DT 07-NOV-2001 (first entry)
DT 07-NOV-2001 (first entry)
XX

PN 07-NOV-2001 (first entry)
XX

Human immune/haematopoietic a
XX
Human; immune; haematopoietic A
XX
Cytostatic; gene therapy; vac
XX

PN W0200157182-A2.
XX

PN W0200157186300

PN W0200157186300

PN W02001571
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2000US-023575
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2000US
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XX
AC AD1125
XX
AC AD1125
XX
DT 22-APR
DT 22-APR
DX Mutant
XX
KW ds; ca
KW ds; ca
KW ovaria
XX
CS Homo 8
XX
PN W02003
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           antino acid sequences given in AAMS170 to AAMS1921. (I) have cytostatic activity, and can be used in gene therapy and vaccine production. (I) concerns and polynucleotides may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate (I) expression. For example, they may be used to treat disorders associated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of (I) by expressing inactive proteins or to supplement the patients own production of (I). Additionally, (I) collections into a host cell and culturing the cell to express the concept and treat immune/haematopoietic may be used to prevent, and cancers and cancer metastases of haematopoietic actived cells. AAK64703 conners and cancer metastases of haematopoietic actived cells. AAK64703 conners from the present invention. AAK54942 to AAK54950 and AAM82169 crepresent sequences used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local S
Matches 55
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17-NOV-2000; 2000US-0249300P.
01-DEC-2000; 2000US-0250160P.
01-DEC-2000; 2000US-0250391P.
05-DEC-2000; 2000US-0251988P.
05-DEC-2000; 2000US-0251479P.
06-DEC-2000; 2000US-0251479P.
08-DEC-2000; 2000US-0251856P.
08-DEC-2000; 2000US-0251868P.
08-DEC-2000; 2000US-0251989P.
08-DEC-2000; 2000US-0251989P.
08-DEC-2000; 2000US-0251989P.
11-DEC-2000; 2000US-0251990P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nucleic
                                                                                                        ds; cancer;
                                                                                                                                        Mutant
                                                                                                                                                                     22-APR-2004
                                                                                                                                                                                                                                ADI12543
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAK54951 to AAK64702 encode the human immune/haematopoietic antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (HUMA-) HUMAN GENOME SCI INC
                                              Homo sapiens
                                                                             ovarian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2001-483426/52
                                                                                                                                                                                                                                                                                                                                         837
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                                                                                                                                                                                                                                                                                                          57
                                                                                                                                                                                                                                                                                                                                                                       55;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              for
                                                                                           cancer susceptibility
                                                                                                                                      human BRCA1 genomic DNA resulting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           acids encoding
                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                             cancer;
                                                                                                                                                                                                                                                                                                                            GATCTGCCTGCCTCGGCCTCCCAAAGTGCTGGGATTACAGGCGTGAGCCACCACCGCC 893
                                                                                                                                                                                                                                                                                                                                                                                                                                     57
                                                                                                                                                                                                                                standard; DNA;
                                                                                                                                                                                                                                                                                                            GATCTGCCTGCCTCAGCCTCCCAAAGTGCTGGGATTACAGGCATGAGCCACCACGCC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Barash SC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            preventing,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQ ID NO 38773; 3071pp + Sequence Listing; English
                                                                                                                                                                                                                                                                                                                                                                                                                                     BP;
                                                                                        human; tumour suppressor; er susceptibility gene 1; BRCA1; repetitive
                                                                                                                                                                                                                                                                                                                                                                      Conservative
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                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                     10 A; 14 C; 21 G; 12 T; 0 U; 0 Other;
                                                                             recombination; mutant
                                                                                                                                                                                                                                                                                                                                                                                      5.4%;
96.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           human immune/hematopoietic diagnosing and/or treating
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                                                                                                                                                                                                                                  ВP
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                                                                                                                                                                                                                                                                                                                                                                       Score 53.8; D
Pred. No. 53;
0; Mismatches
                                                                                                                                        from deletion 3 SeqID
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RESULT 14
ACD93419/c
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Best Local (
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                                                                                                                                                                            Open reading frame detection; genome sequencing; colon cancer; breast cancer; population genome analysis; genetic shift; cancer; antibiotic resistance; antibiotic non-tolerance; congenital disease; agriculture; food crop genome; resistance gene; retrovirus; influenza virus; eukaryotic pathogen detection; trypanosome; Plasmodium;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   present invention describes newly discovered deletion mutations that are believed to be deleterious and cause significant alterations in the structure or biochemical function of BRCA1. Accordingly, it provides methods for detecting such mutants, as well as identifying and screening for cytostatic compounds useful for treating or preventing cancers associated with a BRCA1 genetic variant. This polynucleotide is a mutant human BRCA1 genomic DNA fragment that arrises as a result of a recombination event (deletion 3), which causes the omission of exons 16 and 17, given in an exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Predicting a predisposition to cancer in a patient comprising detecting deletion in the BRCA1 gene that results from the unequal crossover between a pair of repetitive sequences in the BRCA1 gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 07-JUN-2002; 2002US-0387132P
09-AUG-2002; 2002US-0402430P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         09-JUN-2003; 2003WO-US018098
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; SEQ ID NO 26; 59pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2004-062369/06.
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                                                                                                                                                                                                                                                                                                                                                                                   ACD93419 standard; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 60 BP; 14 A; 16 C; 20 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                               Human colon
                                                                                                                                                                                                                                                                                                                 23-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         deletion indicates a predisposition to breast and ovarian cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (MYRI-) MYRIAD GENETICS INC
                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          825 TCTGGACCTTGTGATCTGCCTGCCTCGGCCTCCCAAAGTGCTGGGATTACAGGCGTGAGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     60 TCCTGACCTTGTGATCTGCCCGCCTCGGCCTCCCAAAGTGCTGGGATTACAGGCGTAAGC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         56;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
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                                                                                                                                                                                                                                                                                 cancer cell expressed cDNA #1831.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        5.4%;
93.3%;
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Pred.
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20-NOV-1998; 27-SEP-1999; 24-OCT-2002.

98US-00196716 99US-00406117 JS2002155438-A1

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RESULT 15
ACC84454
ID ACC84
XX
AC ACC84
XX
DT 28-AU
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention describes a method of determining open reading frames in CC the genome of organism, comprising contacting mRNA from cell of organism CC with a single oligonucleotide primer (I) at low stringency, preparing CC single-stranded cDNA by reverse transcribing mRNA with (I), amplifying CC cDNA, sequencing the product, and repeating the contacting, preparing CC and amplifying steps with different primers and sequencing resulting nucleic acids. The method is useful for: determining that a known CC nucleotide sequence from a genome of an organism corresponds to a CC nucleotide sequence from a genome of an organism corresponds to a CC nucleotide sequence of an open reading frame; for preparing a contig, CC all or part of a genome of an organism. and for sequencing CC all or part of subjects and can be used to carry out genetic analyses of CC long cancer or breast cancer cell. The method is useful for analyses of CC populations of subjects and can be used to carry out genetic analyses of CC populations of subjects and can be used to study living CC systems to determine if, e.g. there have been genetic shifts which render CC an individual or population more or less likely to be afflicted with CC diseases such as cancer, to determine antibiotic resistance or non-CC tolerance, and so forth. The method can also be used in the study of CC congenital diseases, and the risk of affliction to a foetus, as well as the study of whether the conditions are likely to be passed to offspring CC carried out in all animals, plants, birds, fish, etc. Using this method, in the area of agriculture, for example the genomes of food crops can be considered to determine if resistance genes are present, defects in plant CC determination of the pathogens which integrate into the genome, such as constant of the pathogens which integrate into the genome, such as constant of the pathogens within the pathogens with a fifteen into the genome, such as constant of the pathogens with the constant of the pathogens within the constant of the pat
                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity Matches 57; Conserv
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(NETO/)
(BREN/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          retroviruses and other integrating viruses such as influenza virus, have undergone shifts or mutations, which may require different approaches to therapy. This method is also applied to the astropens, such as trypanosomes, different types of plasmodium, etc. The method essentially eliminates sequencing of non-coding portions. This sequence represents a polynucleotide isolated from human colon cancer cell cDNA library
                       28-AUG-2003
                                                                      ACC84454;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 9; Page 286; 959pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2003-182626/18
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                                                                                                              ACC84454 standard;
                                                                                                                                                                                                                                                                        750 CCAC 753
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                                                                                                                                                                                                                                                                                                                                               CCTCCCGGGTTCAAGTTATTCTCCTGCCCCAGCCTCCTGAGTAGCTGGGACTACAGGCGC 749
                                                                                                                                                                                                                                                                                                                     CCTCCCGGGTTCATGCCATTCTCCTGCCTCAGCCTCCCGAGTAGCTGGGACTACAGGCAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP; 13
                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                         (first entry)
                                                                                                                DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; 15 C; 27 G; 10 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                         Score 52.8; Di
Pred. No. 66;
O; Mismatches
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RESULT 16
ADI20585
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                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            referred to as cell death peptide. Thought to be cytostatic, antibacterial, immunosuppressive and antiinflammatory. It is useful for treating a condition in a patient requiring removal or destruction of cells, for treating a condition such as benign or malignant tumor, inflammatory disease, autoimmune disease and infectious disease. The peptide useful for treatment is derived from the amino acid sequence for a pancreatic thread protein. The peptide is conjugated, linked or bound to a molecule chosen from antibody or its fragment, antibody-like binding molecule, where the molecule has a higher affinity for binding to a tumor or other target than binding to other cells. Treatment using NTP peptides can remove benign tumors with less risk and fewer of the undesirable side
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19-JUL-2001; 2001US-0306150P.
19-JUL-2001; 2001US-0306161P.
16-NOV-2001; 2001US-0331477P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 can remove benign tumors with research is an NTP encoding sequence effects of surgery. The present sequence is an NTP encoding sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel neural thread protein peptide, referred as cell death peptide, useful for treating prostatic hyperplasia, psoriasis, eczema, dermatos atherosclerosis, cosmetic modification to skin, throat, mouth, muscle.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cytostatic; Antibacterial; Immunosuppressive; Antiinflammatory; neural thread protein; NTP; tumour; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to a neural thread protein (NTP) peptide referred to as cell death peptide. Thought to be cytostatic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Page 15; 77pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NTP peptide encoding sequence #1.
                                              WO2003025229-A1
                                                                            Homo sapiens
                                                                                                                                         Oligonucleotide sequence enquiry #72.
                                                                                                                                                                          15-APR-2004
                                                                                                                                                                                                                                        ADI20585 standard; DNA;
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                                                                                                                                                                                                                                                                                                                                           699 TTCAAGTTATTCTCCTGCCCCAGCCTCCTGAGTAGCTGGGACTACAGGCGC 749
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                                                                                                           ds;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 60 BP; 9 A;
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                                                                                                                                                                          (first entry
                                                                                                                                                                                                                                                                                                                                                                                                5.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                               21 C; 15 G; 15 T; 0 U; 0 Other;
                                                                                                                                                                                                                                         60
                                                                                                                                                                                                                                                                                                                                                                                   ; Score 51; DB
%; Pred. No. 76;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                  DB 1;
                                                                                                                                                                                                                                                                                                                                                                                     0,
                                                                                                                                                                                                                                                                                                                                                                                                                   Length
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RESULT 17
ACC79017/c
ID ACC79
XX ACC79
AC ACC79
AC ACC79
XX Lumar
XX Humar
XX Humar
XX Humar
XX Homo
X
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Best Local S
Matches 54
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention relates to identifying an eRNA or a DNA sequence comprising an eRNA-encoding sequence in the nucleome of a eukaryotic cell comprises identifying non-protein-encoding nucleotide sequences within an mRNA transcript or a DNA sequence encoding same in the nucleome. The methods are useful for identifying an eRNA or DNA for modifying a genetic network in cell to alter the cells phenotype. The present sequence represents human oligonucleotide sequence enquiry.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; regulation; single nucleotide polymorphism; SNP; gene therapy; transcription factor binding site cluster; probe; primer; gene; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ACC79017 standard;
New set of regulatory single nucleotide polymorphism (SNP) polynucleotides, useful in diagnostic assays, in testing s
                                                                                                                                                                                                                                         17-SEP-2001;
30-NOV-2001;
                                                                                                                                                                                                                                                                                                                   11-SEP-2002; 2002WO-US028842
                                                                                                                                                                                                                                                                                                                                                                         27-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                        WO2003025198-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
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                                                                               WPI; 2003-354608/33
                                                                                                                                   Nowotny
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       637
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CTGTCACCCAGGCTGGAGTGCAGTGGCGCCATCATGGCTCACTGCAGCCTCAACCTCCTG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP; 10 A; 22 C; 16 G; 12 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQ ID NO 75; 137pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                       GENOMICS LLC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2001US-0324127P
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                                                                                                                                                                                                                                                              2001US-0322723P
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); Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 50.4;
Pred. No. 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "single nucleotide polymorphism"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention describes a set of regulatory single nucleotide CC polymorphism (SNP) polynucleotides or its complementary set of contiguous CC polynucleotides comprising polynucleotides of at least 6 contiguous CC nucleotides contains a regulatory SNP with 5' and/or 3' genomic flanking sequence. A set of regulatory SNP polynucleotides contains several regulatory SNP swhich collectively map CC polynucleotides contains several regulatory SNP swhich collectively map CC to several transcription factor binding site cluster (TFC) sequences so CC that each SNP lies within a TFC sequence, and a genomic nucleic acid contains a complementary except for the SNP, to a portion of a genomic CC identical or complementary except for the SNP, to a portion of a genomic CC regulatory SNP polynucleotides 3' to the CC regulatory SNP polynucleotides can be used in CC gene therapy. The regulatory SNP polynucleotides can be used in CC regulatory systems for finding new drugs, or for treating or CC polynucleotides are also useful as probes or primers for detecting the regulatory SNPs. The present sequence represents a human genome related CC polynucleotides comprising a SNP, which is used in an example from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 18
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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 2; Page 21; 37pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        finding new drugs, or for treating or preventing a disease associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              with the regulatory SNP.
                                                                                                                                                                                                                                                                                                                                                             ds; cancer; human; tumour suppressor;
breast cancer susceptibility gene 1; BRCA1; repetitive
                                                                                                                                                                                                                                                                                                                                                                                                      Mutant human BRCA1 genomic DNA resulting from
                                                                                                                                                                                                                                                                                                                                                                                                                                    22-APR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           present invention
                                                                                                                                                                                  07-JUN-2002; 2002US-0387132P.
09-AUG-2002; 2002US-0402430P.
                                                                                                                                                                                                                                                          18-DEC-2003
                                                                                                                                                                                                                                                                                                                                              ovarian
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                                                                                                                                                        (MYRI-) MYRIAD GENETICS INC
                                                                                                                                                                                                                            09-JUN-2003; 2003WO-US018098
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                                                                                                   WPI; 2004-062369/06
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CCCAGGCTGGAGTGCAGTGGCGGGATCTCRGCTCACTGCAAGCTCCGCCTCCCAGGTTCA
                                                                                                                              Hendrickson
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Disclosure;

SEQ ID

NO 34; 59pp; English

Predicting a predisposition to cancer in a patient comprising detecting a deletion in the BRCA1 gene that results from the unequal crossover between a pair of repetitive sequences in the BRCA1 gene.

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RRESULT 19
AA179765
ID AA179765
AX AA179765
AX AA179
AX AA179
AX AA179
AX Humar
XX Humar
XX Humar
XX Homo
PN WO201
XX WO201
XX 30-N
PR 30-N
PR 29-N
XX WPI
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PR 107-JI
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PR 107-JI
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Best Local (
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                           AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide sequences (I), which contain single nucleotide polymorphisms (SNPs).

AAM53114 to AAM53329 represent peptides related to human polymorphic polynucleotide sequences. The sequences can be used in gene and protein therapy, and in vaccine production. (I) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of polymorphic polypeptides. For example, (I) may be used to treat disorders by rectifying mutations or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             believed to be deleterious and cause significant alterations in the structure or biochemical function of BRCA1. Accordingly, it provides methods for detecting such mutants, as well as identifying and screening for cycostatic compounds useful for treating or preventing cancers associated with a BRCA1 genetic variant. This polynucleotide is a mutant human BRCA1 genomic DNA fragment that arrises as a polynucleotide in a mutant recombination event (deletion 5), which causes the omission of exons 15 and 16, given in an exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                            Claim 1; Page 2557; 2653pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-356160/37
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29-NOV-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Polymorphic nucleic acid sequences, useful in genetic testing
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Pred. No. 96;
O; Mismatches
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polymorphic polypeptides.
by rectifying mutations of
the activity of polypeptic
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696 GGGTTCAAGTTATTCTCCTGCCCCCAGCCTCCTGAGTAGCTGGGACTACAGGCG

Query Match Best Local

Local

Similarity

4.9%; nilarity 94.3%; Conservative

<u>,</u>

Score 48.2; DB 1; Pred. No. 1.1e+02; 0; Mismatches 3

w ••

Indels

0,

0

Sequence

60 BP; 9 A; 18 C; 18 G; 15 T; 0 U; 0 Other;

represents

human oligonucleotide

sequence

enquiry.

Matches

50;

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RESULT 20
ADI20575
ID ADI20
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        by expressing inactive proteins or to supplement the patients own production of polypeptide. Additionally, (1) and its complementary sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and therefore which patients may be in need of restorative therapy. The polypeptides encoded by (1) may be used as antigens in the production of antibodies specific for polymorphic polypeptides. The antibodies may also be used to down regulate expression and activity. The antibodies may also
                                           The present invention relates to identifying an eRNA or a DNA sequence comprising an eRNA-encoding sequence in the nucleome of a eukaryotic cell comprises identifying non-protein-encoding nucleotide sequences within an mRNA transcript or a DNA sequence encoding same in the nucleome. The methods are useful for identifying an eRNA or DNA for modifying a genetic network in cell to alter the cells phenotype. The present sequence
                                                                                                                                                                                                       Identifying an eRNA or a DNA sequence comprising an eRNA-encoding sequence in the nucleome of a eukaryotic cell, comprising identifying protein-encoding nucleotide sequences within an mRNA transcript or a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Oligonucleotide sequence enquiry #62.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15-APR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADI20575;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADI20575 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            polypeptides in samples
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             be used as diagnostic agents for detecting the presence of polymorphic
                                                                                                                                                            Example 12; SEQ ID NO 65; 137pp; English
                                                                                                                                                                                                                                                                          WPI; 2003-371830/35
                                                                                                                                                                                                                                                                                                                                                                        19-SEP-2001; 2001US-0324127P
                                                                                                                                                                                                                                                                                                                                                                                                        19-SEP-2002; 2002WO-AU001286.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO2003025229-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                                         (UYQU ) UNIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 638
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 49;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ds; eRNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 TGTCGCCCAGGCTGGAGTGCAGTGGCGCAATCTTGGCTCACTGCAACCTC 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TGTCACCCAGGCTGGAGTGCAGTGGCGCAATCTTGGCTCACTGCAACCTC 687
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP; 8 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                         Gagen M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                         QUEENSLAND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             4.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17 C; 15 G; 11 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                          Stanley
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 48.4;
Pred. No. 91;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 7:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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GGGTTCAAGTGATTCTCCTGCCTCAGCCTCCCGAGTAGCTGGGACTACAGGCG 54

16-NOV-2000 AAA77228;

(first entry)

17-NOV-1999;

99WO-US027293 98US-0109024P. 99US-00443199.

25-MAY-2000 WO200029623-A2

17-NOV-1998; 16-NOV-1999;

Shimkets RA, Leach

3

(CURA-) CURAGEN CORP.

WPI; 2000-387826/33

variation

Location/Qualifiers replace(26,C)
/*tag= a

Homo

sapiens

Human; single nucleotide polymorphism; SNP; chromosome 15; detection;

identification; gene therapy; ss.

Human clone cg43971764 polymorphic site, SEQ ID NO:911.

AAA77228 standard; cDNA; 51 BP

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CE silent SNPs. Sequences 1113 to 1192 (AAA77430-A77509) are consecutive CC pairs of nucleotides containing SNPs which result in changes in the CC corresponding maino acid sequences (ABA17439-B11828). The SNPs in CC sequences 1113 to 1128 (AAA77430-A77445) lead to conservative amino acid changes, while those in sequences 1129 to 1186 (AAA77446-A77503) result CC in non- conservative changes. The SNPs in sequences 1187 to 1192 (C (AAA77504-A77509) generate frameshift mutations. The invention also CC relates to a method of detecting a polymorphic site in a nucleic acid and CC a method of detecting polymorphic site in a nucleic acid and CC against such peptides containing polymorphic sites, antibodies raised CC gene therapy of an individual having, suspected of having, or at risk of CC gene therapy of an individual having, suspected of having, or at risk of CC gene therapy of an individual having, suspected of having, or at risk of CC gene therapy of the condition due to the presence of a sequence CC polymorphism. Such treatment would comprise administration of the wild-cype nucleic acid sequence. Antibodies raised against polymorphic CC type nucleic acid sequence. Antibodies raised against polymorphic
밁
                         Ş
                                                      Query Match
Best Local S
Matches 49
                                                                                                               Sequence
                             847
                                                        l Similarity
49; Conserv
                                                                                                                  ភ្ន
                CCTCGGCCTCCCAAAGTGCTGGGATTACAGGCGTGAGCCACCACCACCGCC 897
   CCTCAGCCTCCCAAAGTGCTGGGATTACAGGCATGAGCCACCACGCCCGGC 51
                                                                                                               BP; 11
                                                          Conservative
                                                                                                                 Þ
                                                                      4.8%;
                                                                                                                 20 C; 13 G; 7 T; 0 U; 0 Other;
                                                          0,
                                                                      Score 47.8;
Pred. No. 97;
                                                          Mismatches
                                                                                       8
                                                                                   1; Length 51;
                                                          ب
                                                          Indels
                                                          0
                                                          Gaps
                                                          0
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RESULT 23 AAI78079 ID AAI78 XX

AAI78079 standard; DNA; 51

ВP

0

Sequences AAA76318-A77509 represent 1192 human nucleic acid sequences which contain single nucleotide polymorphisms (SNPs). Sequences 1 to 1112 (AAA76318-A77429) are consecutive pairs of nucleotides which contain

Claim 1; Page 433; 543pp; English. presence of a sequence polymorphism.

Human nucleic acids containing single nucleotide polymorphisms, useful for treating a subject suffering, or at risk from a pathology due to t

to the

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RESULT 22
AAI76251/c
ID AAI762
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            X##X#X#X###X###X##DDDDDDDDDDDDDDDDX&
                                                                                                                                                                       CC Equences (1), which contain single nucleotide polymorphic polymorphic CC AAM53114 to AAM5329 represent peptides related to human polymorphic CC polymucleotide sequences. The sequences can be used in gene and protein CC therapy, and in vaccine production. (I) and the polymeptides encoded by the may be used in the prevention, diagnosis and treatment of diseases cc associated with inappropriate expression of polymorphic polypeptides. For CC deletions in a patient's genome that affect the activity of polypeptides by expressing inactive proteins or to supplement the patients own CC gequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and CC therefore which patients may be used as antigens in the production of polypeptides encoded by (I) may be used as antigens in the production of polypeptides encoded by (I) may be used as antigens in the production of be used to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the presence of polypeptides.
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                           S
                                                              Matches
                                                                                              Query Match
                                                                                Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               quantitation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; single nucleotide polymorphism; SNP; protein therapy; vaccine; probe; diagnostic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human silent SNP containing nucleic acid SEQ:3192
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   09-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAI76251;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Polymorphic nucleic acid sequences, useful in genetic testing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30-NOV-1999; 99US-0168138P.
29-NOV-2000; 2000US-00726173.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200140521-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                               Sequence 51 BP; 12 A; 16 C; 15 G; 8 T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; Page 1027; 2653pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-356160/37.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Shimkets RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30-NOV-2000; 2000WO-US032758.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAI73060 to AAI79867 represent isolated human polymorphic polymucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (CURA-) CURAGEN CORP.
                           639 GTCACCCAGGCTGGAGTGCAGTGGCGCAATCTTGGCTCACTGCAACCTCTG
51
                                                                49;
                                                                                                                                                                as diagnostic agents for tides in samples
                                                                                Similarity
GTCACCCAGGCTGGAGTGCAGTGGCGTGATCTTGGCTCACTGCAACCTCTG 1
                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Leach M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               py; vaccine; probe; diagnostic assay;
restorative therapy; polymorphic; ds.
                                                                              4.8%;
                                                                Score 47.8; D
Pred. No. 97;
0; Mismatches
                                                                0;
                                                                                                                                 U; 0 Other;
                                                                                                DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    genome; gene therapy;
                                                                  2
                                                                                                 Length 51;
                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      detection;
                                                                    0
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                                                                    Gaps
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RESULT 24
AAI73248/c
ID AAI732
XX
AC AAI732
XX
DT 09-NOV
XX
DE Human
XX
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                                                                                                                                                                                                                                                                                                                                                                              CC AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide CC GAM53114 to AAM53129 represent peptides related to human polymorphic (SNPs). CC AAM53114 to AAM53129 represent peptides related to human polymorphic CC polynucleotide sequences. The sequences can be used in gene and protein CC therapy, and in vaccine production. (I) and the polypeptides encoded by CC them may be used in the prevention, diagnosis and treatment of diseases CC associated with inappropriate expression of polymorphic polypeptides. For CC deletions in a patient's genome that affect the activity of polypeptides or CC deletions in a patient's genome that affect the patients own CC production of polypeptide. Additionally, (I) and its complementary CC sequences may also be used as DNA probes in diagnostic assays to detect CC and quantitate the presence of similar nucleic acids in samples, and CC polypeptides encoded by (I) may be used as antigens in the production of colypeptides encoded by (I) may be used as antigens in the production of CC be used to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic.
                                                                                                                                                                                                                                                                   Query Match
Best Local S
Matches 49
               Human silent SNP containing nucleic acid
                                                                                                                                                                                                                                                                                                                                    Sequence 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; Page 2046; 2653pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2001-356160/37.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-NOV-1999; 99US-0168138P.
29-NOV-2000; 2000US-00726173.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               30-NOV-2000; 2000WO-US032758.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; single nucleotide polymorphism; SNP; genome; gene therapy; protein therapy; vaccine; probe; diagnostic assay; detection; quantitation; restorative therapy; polymorphic; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human silent
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAI78079;
                                                  09-NOV-2001
                                                                                                                   AAI73248 standard; DNA; 51 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Polymorphic nucleic acid sequences, useful in genetic testing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (CURA-)
                                                                                                                                                                                                                                  641 CACCCAGGCTGGAGTGCAGTGGCGCAATCTTGGCTCACTGCAACCTCTGCC 691
                                                                                                                                                                                                     ŭ
                                                                                                                                                                                                                                                                   49;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURAGEN CORP.
                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ₽Ą,
                                                                                                                                                                                                     CACCCAGGCTGGAGTGCAGTGGCGCAATCTCGGCTCACTGCAACCTCCGCC 51
                                                                                                                                                                                                                                                                                                                                      BP; 9 A; 20 C; 14 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SNP containing nucleic acid
                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                       in samples
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Leach M;
                                                                                                                                                                                                                                                                                   4.8%;
96.1%;
                                                                                                                                                                                                                                                                   0; Mismatches
                                                                                                                                                                                                                                                                                 Score 47.8;
Pred. No. 97
               SEQ:189
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQ:5020
                                                                                                                                                                                                                                                                                                    DB 1; Length 51;
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                                                                                                                                                                                                                                                                     Indels
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                                                                                                                                                                                                                                                                   Gaps
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RESULT 25
AD112542/c
ID AD1125
XX AD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CC AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide CC sequences (I), which contain single nucleotide polymorphisms (SNPs). CC AAM53114 to AAAM53129 represent peptides related to human polymorphic CC AAM53114 to AAAM53129 represent peptides related to human polymorphic CC polynucleotide sequences. The sequences can be used in gene and protein CC therapy, and in vaccine production. (I) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases CC associated with inappropriate expression of polymorphic polypeptides. For CC example, (I) may be used to treat discorders by rectifying mutations or CC example, (I) may be used to treat discorders by rectifying mutations or CC production in a patient's genome that affect the activity of polypeptides by expressing inactive proteins or to supplement the patients own CC production of polypeptide. Additionally, (I) and its complementary CC sequences may also be used as DNA probes in diagnostic assays to detect CC and quantitate the presence of similar nucleic acids in samples, and CC therefore which patients may be in need of restorative therapy. The CC sequences may also be used as antigens in the production of antibodies may also be used as detection of antibodies may also be used as detection the antibodies may also be used as detection the antibodies may also be used as detection the presence of polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                               ADI12542 standard; DNA; 49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 51 BP; 12 A; 8 C; 23 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Polymorphic nucleic acid sequences, useful in genetic testing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2001-356160/37.
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29-NOV-2000; 2000US-00726173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30-NOV-2000; 2000WO-US032758
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200140521-A2
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                                      Homo sapiens
                                                                                                      ovarian cancer; recombination; mutant.
                                                                                                                                             breast cancer susceptibility
                                                                                                                                                                         ds; cancer; human; tumour suppressor;
                                                                                                                                                                                                                                     Mutant human BRCA1 genomic DNA
                                                                                                                                                                                                                                                                                                                22-APR-2004
                                                                                                                                                                                                                                                                                                                                                                               ADI12542;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 polypeptides in samples
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              be used as diagnostic agents for detecting the presence of polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (CURA-) CURAGEN CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        974
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           49;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Page 113; 2653pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CTCACTGCAACCTCTGCCTCCCAGGCTCAAGCGATTCTCCTGCCTCAGCCT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CTCACTGCAACCTCTGCCTCCCGGGCTCAAGCGATTCTCCCTGTCTCAGCCT 1024
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Leach M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  99US-0168138P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 47.8; D. Pred. No. 97; 0; Mismatches
                                                                                                                                                                                                                                            resulting
                                                                                                                                         1; BRCA1; repetitive Alu,
                                                                                                                                                                                                                                                from deletion 3 SeqID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  멂
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1; Length 51;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           <u>۷</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
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                                                                                                                                                                                                                                                                                                                                                                AAI79764
                                                                                                                                                                                                                                                                                                                                                                                 RESULT 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This invention relates to a novel method for predicting a predisposition to cancer in a patient by detecting large deletions in the human tumour suppressor gene identified as the breast cancer susceptibility gene 1 (BRCA1). Specifically, it refers to deletions that result from the unequal crossover between a pair of repetitive Alu sequences in the BRCA1 gene, such that the recombined nucleotide sequence containing the deletion indicates a predisposition to breast and ovarian cancer. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-JUN-2002; 2002US-0387132P.
09-AUG-2002; 2002US-0402430P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               09-JUN-2003; 2003WO-US018098
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO2003104474-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18-DEC-2003
                                                                                                                                                                          Human; single nucleotide polymorphism; SNP; genome; gene therapy; protein therapy; vaccine; probe; diagnostic assay; detection; quantitation; restorative therapy; polymorphic; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               methods for detecting such mutants, as well as identifying and screening for cytostatic compounds useful for treating or preventing cancers associated with a BRCA1 genetic variant. This polynucleotide is a mutant human BRCA1 genomic DNA fragment that arises as a result of a recombination event (deletion 3), which causes the omission of exons 16 and 17, given in an exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  present invention describes newly discovered deletion mutations that are believed to be deleterious and cause significant alterations in the structure or biochemical function of BRCAL Accordingly, it provides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; SEQ ID NO 25; 59pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            predicting a predisposition to cancer in a patient comprising detecting deletion in the BRCA1 gene that results from the unequal crossover
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Scholl T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 49 BP; 11 A; 13 C; 17 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          between a pair of repetitive sequences in the BRCA1 gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2004-062369/06
30-NOV-1999; 99US-0168138P.
29-NOV-2000; 2000US-00726173.
                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                             Human nonconservative
                                                                                                                                                                                                                                                                                09-NOV-2001
                                                                                                                                                                                                                                                                                                                                                AAI79764 standard; DNA; 51 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (MYRI-) MYRIAD GENETICS INC
                                                                                                               WO200140521-A2
                                                30-NOV-2000; 2000WO-US032758
                                                                                07-JUN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                    49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    48;
                                                                                                                                                                                                                                                                                                                                                                                                                                  ACCTTGTGATCTGCCCGCCTCGGCCTCCCAAAGTGCTGGGATTACAGGC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ACCTTGTGATCTGCCTGCCTCGGCCTCCCAAAGTGCTGGGATTACAGGC 878
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hendrickson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4.8%;
                                                                                                                                                                                                                                             amino acid changing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ward
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 47.4;
Pred. No. 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       99;
                                                                                                                                                                                                                                                SNP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1; Length
                                                                                                                                                                                                                                                nucleic acid SEQ:6705
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         49;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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CC sequences (1), which contain single nucleotide polymorphisms (SNPs).

CC AAM53114 to AAM53329 represent peptides related to human polymorphic CC polymucleotide sequences. The sequences can be used in gene and protein CC therapy, and in vaccine production. (1) and the polymorphise secondary, and in vaccine production, diagnosis and treatment of diseases CC associated with inappropriate expression of polymorphic polymorphic contents or contents in a patient's genome that affect the activity of polypoptides. For CC example, (1) may be used to treat disorders by rectifying mutations or conduction in a patient's genome that affect the activity of polypoptides by expressing inactive proteins or to supplement the patients own CC production of polypoptide. Additionally, (1) and its complementary CC sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and CC therefore which patients may be in need of restorative therapy. The CC antibodies specific for polymorphic polypoptides. The antibodies may also be used as antigens in the production of construction of constructions of construction of constructions of construction of constructions of constructions
                                                                                                                                                                                                                             Query Match
Best Local S
                                                                                                                                                                                      Matches
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                                                                                                                                                                                                                                                                                                                                                                       Sequence 51 BP; 8 A; 16 C; 15 G; 12 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1; Page 2557; 2653pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-356160/37.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAI73060 to AAI79867
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Polymorphic nucleic acid sequences, useful in genetic testing and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    polypeptides in samples
                                            638 TGTCACCCAGGCTGGAGTGCAGTGGCGCAATCTTGGCTCACTGCAACCTC
                                                                                                                                                                                          48;
                                                                                                                                                                                                                                       Similarity
TGTCGCCCAGGCTGGAGTGCAGTGGTGCAATCTTGGCTCACTGCAACCTC
                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                             4.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  represent isolated human polymorphic polynucleotide
                                                                                                                                                                                          0
                                                                                                                                                                                                                                       Score 46.8;
Pred. No. 1.
                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                       .1e+02
                                                                                                                                                                                                                                                                                             DB 1;
                                                                                                                                                                                                                                                                                        Length
                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                             51;
              50
                                                                                                       687
                                                                                                                                                                                               0;
                                                                                                                                                                                          Gaps
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RESULT 27
AAI73066/c
human; single nucleotide polymorphism; SNP; genome; gene therapy;
protein therapy; vaccine; probe; diagnostic assay; detection;
                                                                                                                                                                                                                  09-NOV-2001
                                                                                                  07-JUN-2001.
                                                                                                                                       Homo sapiens
                                                                                                                                                         quantitation;
                                                                                                                                                                   protein therapy; vaccine;
                                                                                                                                                                                             Human silent SNP containing nucleic acid SEQ:7.
                                                                                                                                                                                                                                     AAI73066;
                                                                                                                                                                                                                                                        AAI73066 standard;
                 Shimkets RA,
                                                     30-NOV-1999; 99US-0168138P.
29-NOV-2000; 2000US-00726173.
                                                                               30-NOV-2000; 2000WO-US032758
                                                                                                                    WO200140521-A2
                                   (CURA-)
                                   CURAGEN CORP.
                                                                                                                                                                                                                  (first entry)
                 Leach M;
                                                                                                                                                          restorative
                                                                                                                                                                                                                                                        DNA;
                                                                                                                                                                                                                                                          51
                                                                                                                                                          therapy; polymorphic;
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WPI; 2001-356160/37

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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            by expressing inactive proteins or to supplement the patients own production of polypeptide. Additionally, (I) and its complementary sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and therefore which patients may be in need of restorative therapy. The polypeptides encoded by (I) may be used as antigens in the production of antibodies specific for polymorphic polypeptides. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic be used as diagnostic agents for detecting the presence of polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             therapy, and in vaccine production. (I) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of polymorphic polypeptides. For example, (I) may be used to treat disorders by rectifying mutations or deletions in a patient's genome that affect the activity of polypeptides between in a patient's genome that affect the activity of polypeptides
                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                      Human; single nucleotide polymorphism; SNP; chromosome 15; detection; identification; gene therapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                     Human clone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAA77229
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAA77229 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            sequences (I), which contain single nucleotide polymorphisms (SNPs).
AAM53114 to AAM53329 represent peptides related to human polymorphic
polynucleotide sequences. The sequences can be used in gene and protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 1; Page 56; 2653pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Polymorphic nucleic acid sequences, useful in genetic testing and
                                           WPI; 2000-387826/33.
                                                                                                                                      17-NOV-1998;
16-NOV-1999;
                                                                                                                                                                                     17-NOV-1999;
                                                                                                                                                                                                                                                                                              variation
                                                                                                                                                                                                                                                                                                                                                                                                                                                   16-NOV-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               polypeptides in samples
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAI73060 to AAI79867 represent isolated human polymorphic
                                                                                                                                                                                                                                                WO200029623-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       638
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       48;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                         CURAGEN CORP.
                                                                           RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TGTCACCCAGGCTGGAGTGCAGTGGCGCAATCTTGGCTCACTGCAACCTC 687
                                                                                                                                                                                                                                                                                                                                                                                                                   cg43971764 polymorphic site, SEQ ID NO:912
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP; 10 A; 15 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                         Leach MD
                                                                                                                                      98US-0109024P
99US-00443199
                                                                                                                                                                                     99WO-US027293
                                                                                                                                                                                                                                                                             replace(26,T)
/*tag= a
                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cDNA; 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
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Pred. No. 1.1e
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 1; Length 51; .1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                u; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  For
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Human nucleic acids containing single nucleotide polymorphisms, useful for treating a subject suffering, or at risk from a pathology due to the

AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide

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RESULT 29
AAI73249/c
ID AAI732
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              relates to a method of detecting a polymorphic site in a nucleic acid and a method of determining the relatedness of two nucleic acids. It also encompasses peptides containing polymorphic sites, antibodies raised against such peptides, and a method of detecting polymorphic proteins/peptides using the antibodies. The nucleic acids are useful for gene therapy of an individual having, suspected of having, or at risk of developing a pathological condition due to the presence of a sequence polymorphism. Such treatment would comprise administration of the wild-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequences AAA76318-A77509 represent 1192 human nucleic acid sequences which contain single nucleotide polymorphisms (SNPs). Sequences 1 to 1 (AAA76318-A77499) are consecutive pairs of nucleotides which contain silent SNPs. Sequences 1113 to 1192 (AAA77430-A77509) are consecutive pairs of nucleotides containing SNPs which result in changes in the corresponding amino acid sequences (AAB11749-B11828). The SNPs in sequences 1113 to 1128 (AAA77430-A77445) lead to conservative amino ac
                                                                                                                                                                                                                                                                                                                                                                                                                 Human; single nucleotide polymorphism; SNP; genome; gene therapy; protein therapy; vaccine; probe; diagnostic assay; detection; quantitation; restorative therapy; polymorphic; ds.
                                                                                                                                                                                                                        30-NOV-1999;
29-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       09-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAI73249 standard; DNA; 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 51 BP; 11 A; 21 C; 13 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        changes, while those in sequences 1129 to 1186 (AAA77446-A77503) result in non- conservative changes. The SNPs in sequences 1187 to 1192 (AAA77504-A77509) generate frameshift mutations. The invention also
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   presence of a sequence polymorphism
                              Claim 1; Page 113; 2653pp;
                                                                                  Polymorphic nucleic acid sequences,
                                                                                                                    WPI; 2001-356160/37
                                                                                                                                                      Shimkets
                                                                                                                                                                                                                                                                          30-NOV-2000; 2000WO-US032758
                                                                                                                                                                                                                                                                                                              07-JUN-2001.
                                                                                                                                                                                                                                                                                                                                              WO200140521-A2
                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human silent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     type nucleic
                                                                                                                                                                                        (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             847 CCTCGGCCTCCCAAAGTGCTGGGATTACAGGCGTGAGCCACCACGCCCGGC 897
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 48;
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                                                                                                                                                      ₽Ą,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CCTCAGCCTCCCAAAGTGCTGGGATCACAGGCATGAGCCACCACGCCGGC 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    leic acid sequence. Antibodies raised against polymorphic can also be used in the treatment of such individuals
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                        99US-0168138P.
2000US-00726173.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SNP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                      Leach
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 containing nucleic acid SEQ:190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4.7%;
                                English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 46.2; DB 1;
Pred. No. 1.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                     useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1;
                                                                                     'n
                                                                                  genetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 51;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                conservative amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                    testing
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RESULT 30
AAI78078
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CC AAM53114 to AAM53329 represent peptides related to human polymorphic polymucleotide sequences. The sequences can be used in gene and protein CC therapy, and in vaccine production. (I) and the polymorphic server, and in vaccine production, diagnosis and treatment of diseases CC associated with inappropriate expression of polymorphic polymoptides. For CC example, (I) may be used to treat disorders by rectifying mutations or CC example, (I) may be used to treat disorders by rectifying mutations or CC example, (I) may be used to supplement the patients own CC polymorphic proteins or to supplement the patients own CC production of polypeptide. Additionally, (I) and its complementary CC sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and CC polypeptides encoded by (I) may be used of restorative therapy. The CC polypeptides encoded by (I) may be used as antigens in the production of CC polypeptides encoded by (I) may be used as antigens in the production of Detect and as propertides. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic polymorphic polymorphic polymorphic agents for detecting the presence of polymorphic p
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; single nucleotide polymorphism; SNP; genome; gene therapy; protein therapy; vaccine; probe; diagnostic assay; detection; quantitation; restorative therapy; polymorphic; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAI78078 standard; DNA;
AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide sequences (I), which contain single nucleotide polymorphisms (SNPs). AAM53114 to AAM53329 represent peptides related to human polymorphic polynucleotide sequences. The sequences can be used in gene and protein therapy, and in vaccine production. (I) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human silent SNP containing nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    09-NOV-2001
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29-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30-NOV-2000; 2000WO-US032758
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          07-JUN-2001
                                                                                                                                                                                                                                                                         Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                        Polymorphic nucleic acid sequences,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2001-356160/37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200140521-A2
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                                                                                                                                                                                                                                                                         Page
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                     2046; 2653pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Leach M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               99US-0168138P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                51
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Pred.
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No. 1
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                                                                                                                                                                                                                                                                                                                                                                                        genetic testing
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associated with inappropriate expression

of polymorphic

polypeptides.

sequences (1), which contain single nucleotide polymorphisms (SNPs).

AAM53114 to AAM53329 represent peptides related to human polymorphic polymucleotide sequences. The sequences can be used in gene and protein therapy, and in vaccine production. (I) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of polymorphic polypeptides. For example, (I) may be used to treat disorders by rectifying mutations or deletions in a patient's genome that affect the activity of polypeptides by expressing inactive proteins or to supplement the patients own production of polypeptide. Additionally, (I) and its complementary sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and

For

AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide

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RESULT 31
AAI79818
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Query Match
Best Local S
Matches 48
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human, single nucleotide polymorphism; SNP; genome; gene therapy; protein therapy; vaccine; probe; diagnostic assay; detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human nonconservative amino acid changing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                            30-NOV-1999; 99US-0168138P
29-NOV-2000; 2000US-00726173
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAI79818;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAI79818 standard;
                                                                                                                                                                               Claim 1; Page 2573; 2653pp; English
                                                                                                                                                                                                                 Polymorphic nucleic acid sequences, useful
                                                                                                                                                                                                                                               WPI; 2001-356160/37
                                                                                                                                                                                                                                                                        Shimkets RA,
                                                                                                                                                                                                                                                                                                                                                                 30-NOV-2000; 2000WO-US032758
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                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         quantitation;
                                                                                                                                                                                                                                                                                                  (CURA-) CURAGEN CORP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         yy; vaccine; probe; diagnostic ass;
restorative therapy; polymorphic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       entry)
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Pred. No. 1
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                                                                                                                                                                                                                     genetic
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RESULT 32
AA176250/c
ID AA176250/c
XX AA176250/c
XX AA176
XX AA176
XX Huma
XX 
                   CC AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide CC sequences (I), which contain single nucleotide polymorphisms (SNPs). CC AAM53114 to AAM53129 represent peptides related to human polymorphic CC polynucleotide sequences. The sequences can be used in gene and protein CC thermapy, and in vaccine production. (I) and the polypeptides encoded by C them may be used in the prevention, diagnosis and treatment of diseases CC associated with inappropriate expression of polymorphic polypeptides. For CC deletions in a patient's genome that affect the activity of polypeptides or CC deletions in a patient's genome that affect the patients own CC by expressing inactive proteins or to supplement the patients own CC sequences may also be used as NAM probes in diagnostic assays to detect CC and quantitate the presence of similar nucleic acids in samples, and CC therefore which patients may be in need of restorative therapy. The CC service which patients may be in need of restorative therapy. The CC contibodies medically (I) may be used as antigens in the production of CC polypeptides encoded by (I) may be used as antigens in the production of CC contibodies may be in need of restorative therapy. The CC contibodies may also be used as diagnostic agents for detecting the presence of polymorphic continues of polymorphic and activity. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Polymorphic nucleic acid sequences, useful in genetic testing and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2001-356160/37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      30-NOV-1999;
29-NOV-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; single nucleotide polymorphism; SNP; genome; gene therapy; protein therapy; vaccine; probe; diagnostic assay; detection; quantitation; restorative therapy; polymorphic; ds.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 51 BP; 9 A; 16 C; 14 G; 12 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         48;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ŗ,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CGGGTTCAAGCGATTCTCCTGCCTCAGCCTCCTGAGTAGCTGGGACTACAG 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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2000US-00726173.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
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Pred. No. 1.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
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                                                                                AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide sequences (I), which contain single nucleotide polymorphisms (SNPB).

AAM53114 to AAM5329 represent peptides related to human polymorphic polymorphic polynucleotide sequences. The sequences can be used in gene and protein therapy, and in vaccine production. (I) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of polymorphic polypeptides. For example, (I) may be used to treat disorders by rectifying mutations or deletions in a patient's genome that affect the activity of polypeptides by expressing inactive proteins or to supplement the patients own production of polypeptide. Additionally, (I) and its complementary sequences may also be used as DAB probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and therefore which patients may be used as antigens in the production of polypeptides encoded by (I) may be used as antigens in the production of be used to down regulate expression and activity. The antibodies may also be used as diagnostic aspants for detecting the presence of polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; single nucleotide polymorphism; SNP; genome; protein therapy; vaccine; probe; diagnostic assay; quantitation; restorative therapy; polymorphic; ds.
                                                            polypeptides in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; Page 1923; 2653pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Polymorphic nucleic acid sequences, useful in genetic testing and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2001-356160/37
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29-NOV-2000; 2000US-00726173.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human silent SNP containing nucleic acid SEQ:4617.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Leach M;
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Pred. No. 1
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detection;
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Query Match Best Local :

Similarity

4.7%;

Score 46.2; DB 1 Pred. No. 1.2e+02

DB 1;

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Conservative

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Sequence

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                                                                            Query Match
Best Local :
                                                                                                                                                     This invention relates to a novel isolated protein which comprises a human mannosyl transferase having the same sequence as the fully defined 611- or 255-amino acid sequence or its fragment. The invention may be useful for the production of compounds with an antimanic or antidepressant activity whilst the disclosed sequences may be used for gene therapy. The invention also provides a human mannosyl transferase fusion protein and a chromosome 9 fusion protein, both of which result from a chromosome 11 translocation. The human mannosyl transferase and the fusion proteins are useful for diagnosing or predicting the susceptibility to a bipolar disorder and for identifying a compound that modulates the activity of a mannosyl transferase. The present sequence is that of a region of human DNA surrounding a single nucleotide polymorphism within the gene which encodes the human mannosyl transferase of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        06-MAY-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     variation
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     human; mannosyl transferase; antimanic; antidepressant; gene therapy; fusion protein; chromosome 9 fusion protein; chromosome 11 translocation; bipolar disorder; single nucleotide polymorphism; SNP; ds.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    New polypeptide comprising human mannosyl transferase, useful for diagnosing or predicting the susceptibility to a bipolar disorder identifying a compound that modulates the activity of a mannosyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      02-AUG-2002; 2002WO-US024490
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                                                                                                                                                                                                                                                                                                                                                                                                        Claim 11; SEQ ID NO 59; 147pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2003-268116/26.
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                                                                                                                            Sequence 51 BP; 7 A;
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                                                                            Similarity
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                         GCCTCCCGGGTTCAAGTTATTCTCCTGCCCCAGCCTCCTGAGTAGCTGGGA 739
 GCCTCCCGGGTTCAAGCGATTCTCCTGCCTCAGCCTCCTGAGTAGCTGGGA 51
                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       transferase-related SNP region DNA SeqID59.
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/*tag= a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA;
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                                                             Score 46.2; Di
Pred. No. 1.2e
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This invention relates to novel tissue specific genes and gene clusters cand the proteins encoded by them. The invention includes the 1922 immune CC gene complex, such as TMD0024 (XM 0 60945), TMD1779 (XM 0 60946), TMD0884 (CC (XM 0 60946), TMD00825 (XM 0 60946), TMD1781 (XM 0 80942), TMD0304 (XM 0 60956), TMD0888 (XM 0 89422), TMD0781 (XM 0 60959), bone marrow specific genes consisting essentially of XM 0 62147 (CC and XM 0 61676 or their proteins, kidney specific genes TMD0049 (XM 0 62147 (CC and XM 0 61676 or their proteins, kidney specific genes TMD0039 (XM 0 87351), TMD0190 (XM 0 87157), TMD0242 (XM 0 88369), TMD0335 (XM 0 89960), CC TMD0371, TMD0374, TMD0469 (XM 0 88369), TMD0733, TMD0374, TMD0466 (XM 0 88369), TMD0733, TMD01114 (XM 0 89960), CC 19841) and TMD1148 (XM 0 87108), pancreas specific genes consisting cessentially of TMD0285, TMD0275, TMD0275, TMD0277, TMD0233, TMD0275, TMD0275, TMD0275, TMD0277, TMD0233, TMD0233, TMD0258, TMD0258, TMD0274, TMD0677, TMD0233, TMD0258, TMD0258, TMD0274, TMD0677, TMD0273, TMD0733, TMD0111, and TMD1127 or its encoded polypeptide or the 11q12.2 spleen gene complex, comprising CC TMD0621 (XM 1 66853), TMD10793 (XM 1 66854), TMD1028 (XM 1 66855) and CC TMD0621 (XM 1 66853), TMD01028 (XM 1 66854), TMD1028 (XM 1 66855) and CC CMD0011ds with an immunostimulant, cytostatic, osteopathic, nephrotropic, antidiabetic, ophthalmological or antianaemic activity. The sequences of
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24-APR-2002; 2002US-0374823P.
01-MAY-2002; 2002US-0376558BP.
20-MAY-2002; 2002US-0381366P.
16-AUG-2002; 2002US-041882P.
20-SEP-2002; 2002US-0424336P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1922 immune gene complex; bone marrow specific gene; kidney specific gene; pancreas specific gene; retinal specific gene; liq12 spleen gene complex; immunostimulant; cytostatic; osteopathic; nephrotropic; antidiabetic; ophthalmological; antianaemic; gene thera immune system cell; kidney cell; pancreas cell; retinal cell; spleen cell; reticuloendothelial cell; cell macuration; lymphoid immune system cell; non-lymphoid immune system cell; pancreas; cancer; multiple myeloma; renal failure; glomerular disease; diabetes; retinal degeneration; optic neuritis; glaucoma; anaemia; human; TMD0621; ds.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Composition comprising tissue-specific (e.g. bone marrow, kidney, pancreas or spleen) gene and gene clusters, useful as molecular markers,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2003-854099/79.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  tissue specific gene; tissue specific gene cluster;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 155; SEQ ID NO 211; 358pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    drug targets, or for diagnosing, preventing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (ORIG-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       or treating e.g. neutropenia
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               gene therapy;
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RESULT 36
AAI73067/c
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AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide sequences (I), which contain single nucleotide polymorphisms (SNPs).

AAM53114 to AAM53329 represent peptides related to human polymorphic polymucleotide sequences. The sequences can be used in gene and protein therapy, and in vaccine production. (I) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of polymorphic polypeptides. For example, (I) may be used to treat disorders by rectifying mutations or deletions in a patient's genome that affect the activity of polypeptides by expressing inactive proteins or to supplement the patients own production of polypeptide. Additionally, (I) and its complementary sequences may also be used as DNA probes in diagnostic assays to detect
                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-356160/37
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29-NOV-2000; 2000US-00726173.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Polymorphic nucleic acid sequences, useful in genetic testing and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (CURA-) CURAGEN CORP.
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                                                                                                                                                                                                                                                                                                                                                                                                                 56; 2653pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Leach
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); Mismatches
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Pred. No. 1.
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RESULT 37
AA174554/c
ID AA1745
XX AA1745
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XX Human
XX Human,
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                             CC AAI73366 to AAI79867 represent isolated human polymorphic polynucleotide CC sequences (I), which contain single nucleotide polymorphisms (SNPs). CC AAM53114 to AAM53329 represent peptides related to human polymorphic CC polynucleotide sequences. The sequences can be used in gene and protein CC therapy, and in vaccine production. (I) and the polypeptides encoded by CC them may be used in the prevention, diagnosis and treatment of diseases CC associated with inappropriate expression of polymorphic polypeptides. For CC example, (I) may be used to treat disproders by rectifying mutations or CC example, (I) may be used to treat disproders by rectifying mutations or CC production in a patient's genome that affect the activity of polypeptides CC by expressing inactive proteins or to supplement the patients own CC production of polypeptide. Additionally, (I) and its complementary CC sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and CC polypeptides encoded by (I) may be used as antigens in the production of antibodies may be in need of restorative therapy. The CC polypeptides encoded by (I) may be used as antigens in the production of antibodies may also be used to down regulate expression and activity. The antibodies may also be used to down regulate expression and activity. The antibodies may also
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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29-NOV-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Polymorphic nucleic acid sequences, useful in genetic testing
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              al Similarity
47; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ₽,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  99US-0168138P.
2000US-00726173.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Leach M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4.6%;
94.0%;
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Pred. No. 1.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
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detection;
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diagnostic agents for detecting the

presence of

polymorphic

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RESULT 38
AAI73064/c
ID AAI730
SXS
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                                        CC AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide CC sequences (I), which contain single nucleotide polymorphisms (SNPs). CC AAM53114 to AAM5329 represent peptides related to human polymorphic CC polynucleotide sequences. The sequences can be used in gene and protein therapy, and in vaccine production. (I) and the polypeptides encoded by CC them may be used in the prevention, diagnosis and treatment of diseases CC associated with inappropriate expression of polymorphic polypeptides. For CC deletions in a patient's genome that affect the activity of polypeptides or CC deletions in a patient's genome that affect the patients own CC by expressing inactive proteins or to supplement the patients own CC sequences may also be used as MAN probes in diagnostic assays to detect CC and quantitate the presence of similar nucleic acids in samples, and CC therefore which patients may be in need of restorative therapy. The CC antibodies mapcific for polymorphic polypeptides. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic polypeptides may also be used as diagnostic agents for detecting the presence of polymorphic polypeptides. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; single nucleotide polymorphism; SNP; genome; gene therapy;
protein therapy; vaccine; probe; diagnostic assay; detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human silent SNP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               polypeptides in
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29-NOV-2000; 2000US-00726173.
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                                                                                                                                                                                                                                                                                                                                                 Claim 1; Page 55; 2653pp; English
                                                                                                                                                                                                                                                                                                                                                                                                   Polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-356160/37
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Shimkets RA, Leach M;
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 Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CTCTGTCACCCAGGCTGGAGTGCAGTGGCGCAATCTTGGCTCACTGCAAC 684
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CTCTGTCACCCAGGCTGGAGTGCAGTGGCACGATCTCGGCTCACTGCAAC 1
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 51
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 BP;
                                                                                                                                                                                                                                                                                                                                                                                                   nucleic acid sequences, useful in genetic testing
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   14
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 10 C;
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Pred. No. 1.
   19 G;
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   8 T; 0
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   U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length
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Query Match Best Local Similarity

4.5%; 95.8%;

Score 44.8; Pred. No. 1

.8; DB 1; . 1.4e+02;

Length 51;

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ARBSULT 39
AAI77324
ID 7AAI77
XX AAI77
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OX H
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                                                                                                                                              Query Match
Best Local
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(2)
                                                                                                                                                                                                                                                                                   by expressing inactive proteins or to supplement the patients own production of polypeptide. Additionally, (I) and its complementary sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and therefore which patients may be in need of restorative therapy. The polypeptides encoded by (I) may be used as antigens in the production of antibodies specific for polymorphic polypeptides. The antibodies may also be used to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic polypeptides in samples.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         polynucleotide sequences. The sequences can be used in gene and protein therapy, and in vaccine production. (I) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of polymorphic polypeptides. For example, (I) may be used to treat disorders by rectifying mutations or deletions in a patient's genome that affect the activity of polypeptides by a procession of polymorphic polypeptides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30-NOV-1999; 99US-0168138P
29-NOV-2000; 2000US-00726173
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide sequences (I), which contain single nucleotide polymorphisms (SNPs). AAM53114 to AAM53329 represent peptides related to human polymorphic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; Page 1815; 2653pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Polymorphic nucleic acid sequences, useful in genetic testing
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                         Sequence 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (CURA-) CURAGEN CORP
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                                                            843 CCTGCCTCGGCCTCCCAAAGTGCTGGGATTACAGGCGTGAGCCACCAC
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                                                                                                                                                       Similarity
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                                                                                                                                                                                                                                         BP; 8 A;
                                                                                                                        Conservative
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                                                                                                                                                 4.5%;
95.8%;
                                                                                                                                                                                                                                         21 C; 14 G; 8 T; 0 U; 0 Other;
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                                                                                                                                                    Score 44.8;
Pred. No. 1
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                                                                                                                                                          .4e+02;
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detection;
                                                                                                                              Indels
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RESULT 40
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                                                                                                                                                                              CC which contain single nucleotide polymorphisms (SNPs). Sequences 1 to 1112 (C (AAA76318 A77429) are consecutive pairs of nucleotides which contain contain contain to 112 (AAA77630 A77509) are consecutive pairs of nucleotides which contain corresponding amino acid sequences (AAA17430-A77509) are consecutive corresponding amino acid sequences (AAA17430-B1828). The SNPs in CC corresponding amino acid sequences (AAA17430-B1828). The SNPs in CC corresponding amino acid sequences 1129 to 1136 (AAA77446-A77503) result conservative changes. The SNPs in sequences 1187 to 1192 (C (AAA77504-A77503) generate frameshift mutations. The invention also CC changes to a method of detecting a polymorphic site in a nucleic acid and CC a method of determining the relatedness of two nucleic acids. It also CC correspondess peptides containing polymorphic site in a nucleic acid and CC against such peptides containing polymorphic sites, antibodies raised CC gene therapy of an individual having, suspected of having, or at risk of C developing a pathological condition due to the presence of a sequence CC type nucleic acid sequence. Antibodies raised against polymorphic wild-contain acid sequence. Antibodies raised against polymorphic
                                                                  Matches
                                                                                Query Match
Best Local :
                                                                                                                                   Sequence 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequences AAA76318-A77509 represent 1192 human nucleic acid sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; Page 515; 543pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human nucleic acids containing single nucleotide polymorphisms, useful for treating a subject suffering, or at risk from a pathology due to the presence of a sequence polymorphism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17-NOV-1998;
16-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25-MAY-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO200029623-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          variation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; single nucleotide polymorphism; SNP; detection; identification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human Alu subfamily SQ gene polymorphic site, SEQ ID NO:1181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          16-NOV-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAA77498;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAA77498 standard; cDNA; 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2000-387826/33.
DB; AAB11817.
                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RΑ,
                          CTGAGCTCAAGCAGTCCACCTGCCTCAGCCTCCCAAAGTGCTGGGATTACA 405
                                                                                                                                                                     can also be used in the treatment of such individuals
 CTGACCTCAAGTGATCCACCTGCCTCAGCCTCCCAAAGTGCTGGGATTACA 51
                                                                                                                                   BP; 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Leach MD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   98US-0109024P.
99US-00443199.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      99WO-US027293
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers replace(26,T) /*tag= a
                                                                                                                                   A; 18 C; 10 G; 11 T; 0 U; 0 Other;
                                                                               4.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ВP
                                                                 0,
                                                                 Score 44.6; D
Pred. No. 1.4e
0; Mismatches
                                                                                  .4e+02;
                                                                                                    ВG
                                                                                                  1,
                                                                  4.
                                                                                                  Length 51;
                                                                  Indels
                                                                  <u>.</u>
                                                                 Gaps
                                                                  0
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밁 S

Query Match Best Local S Matches 47

Local Similarity hes 47; Conserv

4.5%; llarity 92.2%; Conservative

<u>,</u>

Score 44.6; D Pred. No. 1.4e O; Mismatches

1.4e+02; B 1; 4

Length 51; Indels

0

Gaps

0

177

TTAGTAGAGATGGAGTTTCTCCATGTTGGTCAGGCTGGTCTCGAACTCCCG 227

TTAGTAGAGACGGGGTTTCACCATGTTGGTCAGGCTGGTCTCGAACTCCTG 51

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RESULT 41
AAA77230
CC Sequences AAA76318-A77509 represent 1192 human nucleic acid sequences CC which contain single nucleotide polymorphisms (SNPs). Sequences 1 to 1112 CC (AAA76318-A77429) are consecutive pairs of nucleotides which contain (CC silent SNPs. Sequences 1113 to 1192 (AAA77430-A77509) are consecutive pairs of nucleotides containing SNPs which result in changes in the CC corresponding amino acid sequences (AAB11749-B11828). The SNPs in CC sequences 1113 to 1128 (AAA77430-A77445) lead to conservative amino acid CC changes, while those in sequences 1129 to 1186 (AAA77446-A77503) result (CC in non- conservative changes. The SNPs in sequences 1187 to 1192 (CC (AAA77504-A77509)) generate frameshift mutations. The invention also CC entres to a method of detecting a polymorphic site in a nucleic acid and CC against such peptides containing the relatedness of two nucleic acids. It also CC gene therapy of an individual having, suspected of having, or at risk of CC developing a pathological condition due to the presence of a sequence CC polymorphism. Such treatment would comprise administration of the wild-CC peptides can also be used in the treatment of such individuals
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAA77230 standard; cDNA; 51 BP
Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                             Human nucleic acids containing single nucleotide polymorphisms, useful for treating a subject suffering, or at risk from a pathology due to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Shimkets RA, Leach MD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17-NOV-1998;
16-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; single nucleotide polymorphism; SNP; chromosome 8; detection; identification; gene therapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human clone cg43972482 polymorphic site, SEQ ID NO:913
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16-NOV-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAA77230;
                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; Page 433; 543pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                             for treating a subject suffering, or presence of a sequence polymorphism.
                                                                                                                                                                                                                                                                                                                                                                                                                               presence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2000-387826/33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         17-NOV-1999;
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51 BP; 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    98US-0109024P.
99US-00443199.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         99WO-US027293
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         replace(26,C)
/*tag= a
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A; 11 C;
16 G; 15 T; 0 U; 0 Other;
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RESULT 42
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멍
                                                                                                             Query Match
Best Local Similarity
Matches 47; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to oligonucleotides encoding polymorphic variants of proteins related to amylases, amyloid proteins, angiopoietin, apoptosis related proteins, cadherin, cyclin, polymerase, oncogenes, histones, kinases, colony stimulating factors, complement related proteins, cytochromes, kinesins, cytokines, interferons, interleukins, G-protein coupled receptors and thioesterases. The present sequence is one put of a complement of a companion of the peptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of the proteins listed above. Discrete that may be prevented, diagnosed and/or treated includes that may be prevented, diagnosed and/or treated includes the proteins of the proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Shimkets RA, Leach
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     05-JUL-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo
                                                                                                                                                                                                                                                                                                                                      multifactorial diseases with a genetic component, such as autoimmune diseases (e.g. rheumatoid arthritis, multiple sclerosis, diabetes, systemic lupus erythromatosus and Grave's disease), inflammation, cancer (e.g. cancers of the bladder, brain, breast, colon and kidney,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              polymorphic nucleic acids encoding e.g. amylases, cyclins, polymerases, oncogenes and histones, useful for diagnosing and treating, e.g. cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-465210/50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    28-DEC-1999; 99US-0173419P.
27-DEC-2000; 2000US-00173419.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          28-DEC-2000; 2000WO-US035498
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        nervous system disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; Page 2729; 4143pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      autoimmune diseases and infections.
                                                                                                                                                                                                                           Sequence 51
                                                                                                                                                                                                                                                                                                           leukaemia), diseases of the nervous system and an infection of pathogenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sapiens.
                                                        843
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                                CCTGCCTCGGCCTCCCAAAGTGCTGGGATTACAGGCGTGAGCCACCACGCC 893
  CCCGCCTTGGCCTCCCAAAGTGCTGAGATTACAGGCATGAGCCACCACGCC
                                                                                                                                                                                                                              B₽;
                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                              ä
                                                                                                                                                                                                                           A; 20 C; 12 G; 8 T; 0 U; 0 Other;
                                                                                                                                        92.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             51
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                                                                                                                0,
                                                                                                                                           Score 44.6; DB 1
Pred. No. 1.4e+02
                                                                                                                Mismatches
                                                                                                                                                                      DB 1;
                                                                                                                                                                      Length
                                                                                                                   Indels
                                                                                                                                                                         51;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    e.g. cancer,
                                                                                                                0,
        51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cancer;
                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  factor;
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RESULT 43
AAL29843
ID AAL29

AAL29843 standard; DNA; 51

ВP

AAI73062

AAI73062 standard; DNA;

51 ΒP

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RESULT 44
AAI73062/c
ID AAI73'
XX
AC A'
XX
                                                                                                                                                                                                                                                                                     The present invention relates to oligonucleotides encoding polymorphic cc variants of proteins related to amylases, amyloid proteins, angiopoietin, cc apoptosis related proteins, cadherin, cyclin, polymerase, oncogenes, cc proteins, cytochromes, kinesins, cytokines, interferons, interleukins, cg-cc proteins, cytochromes, kinesins, cytokines, interferons, interleukins, G-cc protein coupled receptors and thioesterases. The present sequence is one cc such oligonucleotide. The oligonucleotides and the peptides encoded by chem may be used in the prevention, diagnosis and treatment of diseases cc associated with inappropriate expression of the proteins listed above. Cc Disorders that may be prevented, diagnosis and treatment of diseases cc diseases (e.g. rheumatoid arthritis, multiple sclerosis, diabetes, cc systemic lupus erythromatosus and Grave's disease), inflammation, cancer (e.g. cancers of the bladder, brain, breast, colon and kidney, crossisms) diseases of the nervous system and an infection of pathogenic corpanisms
                                                                                                                                              밁
                                                                                                                                                                           S
                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Immunosuppressive; immunostimulatory; antiinflammatory; cytostatic; neuroprotective; antiinficrobial; gene therapy; vaccine; amylase; cancer; amyloid protein; angiopoletin; apoptosis related protein; cadherin; cyclin; polymerase; oncogene; histone; kinase; colony stimulating factor; complement related protein; cytochrome; kinase; cytokine; interferon; interleukin; G-protein coupled receptor; thioesterase; inflammation; interferon; interleukin; G-protein coupled receptor; thioesterase; inflammation; multifactorial disease; autoimmune disease; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Polymorphic nucleic acids encoding e.g. amylau oncogenes and histones, useful for diagnosing autoimmune diseases and infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human SNP oligonucleotide #3051.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  24-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAL29843;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO200147944-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      nervous system disease; ss.
                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Shimkets RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 27-DEC-2000; 2000US-00173419
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28-DEC-2000; 2000WO-US035498.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens.
                                                                                                                                                                                                                                                                                                                     organisms
                                                                                                                                                                                                                                  Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2001-465210/50.
                                                                                                                                                                   974 CTCACTGCAACCTCTGCCTCCCGGGCTCAAGCGATTCTCCTGTCTCAGCCT 1024
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1; Page 2260; 4143pp; English
                                                                                                                                                 _
                                                                                                                                                                                                                l Similarity 47; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURAGEN CORP.
                                                                                                                                                                                                                                                                                     51
                                                                                                                                                CTCACTGCAGCCTCCACCTCCCGGGCTCAAGCGATTCTCCTGCCTCAGCCT
                                                                                                                                                                                                                                                                                     BP; 7
                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Leach M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   99US-0173419P
                                                                                                                                                                                                                                                                                     A;
                                                                                                                                                                                                                                  4.5%;
92.2%;
                                                                                                                                                                                                                                                                                     24 C; 9 G; 11
                                                                                                                                                                                                                  ٥,
                                                                                                                                                                                                                  Score 44.6; D
Pred. No. 1.4e
0; Mismatches
                                                                                                                                                                                                                                                                                       T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           amylases, cyclins, polymerases,
nosing and treating, e.g. cancen
                                                                                                                                                                                                                                    .4e+02;
                                                                                                                                                                                                                                                      DB
                                                                                                                                                                                                                                                    Length
                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                      <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           e.g. cancer,
                                                                                                                                                                                                                        Gaps
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AAI77522/c
ID AAI777
XX AAI77
XX AAI77
XX AAI77
XX AAI77
XX AAI77
XX AAI77
XX Humar
XX Humar
XX Humar
XW Prote
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CC sequences (I), which contain single nucleotide polymorphisms (SNPs).

CC AAM53314 to AAM53329 represent peptides related to human polymorphic polym
                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Human; single nucleotide polymorphism; SNP; genome; gene therapy; protein therapy; vaccine; probe; diagnostic assay; detection;
                                                                    Human silent SNP containing nucleic acid SEQ:4463.
                                                                                                                                                                                                                7522/c
AAI77522 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-356160/37.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Shimkets RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30-NOV-1999; 99US-0168138P.
29-NOV-2000; 2000US-00726173.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               30-NOV-2000; 2000WO-US032758
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              07-JUN-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; single nucleotide polymorphism; SNP; genome; gene therapy; protein therapy; vaccine; probe; diagnostic assay; detection; quantitation; restorative therapy; polymorphic; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human silent
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                                                                                                                     09-NOV-2001
                                                                                                                                                                    AAI77522;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    polypeptides in samples
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                              47;
                                                                                                                                                                                                                                                                                                                                                                               971
                                                                                                                                                                                                                                                                                                                                   51
                                                                                                                                                                                                                                                                                                                                                           CGGCTCACTGCAACCTCTGCCTCCCGGGCTCAAGCGATTCTCCTGTCTCAG 1021
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                to AAI79867 represent isolated human polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Page 55; 2653pp; English.
                                                                                                                                                                                                                                                                                                                                   CGGCTCACTGCAACCTCCGCCTCCTGGGTTCAAGCGATTCTCCTGCCTCAG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 12 A; 11 C; 21 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   nucleic acid sequences, useful in genetic testing
                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SNP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Leach M;
                                                                                                                     (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  containing nucleic acid SEQ:3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                entry)
                                                                                                                   entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                 4.5%;
                                                                                                                                                                                                                51
                                                                                                                                                                                                                ВP
                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 44.6; DB 1; Length 51; Pred. No. 1.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                           4;
                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
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RESULT 46
AAI78388
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CC sequences (I) may be used as DNA probes in diagnostic asmyles of polypeptides encoded by contains of proteins of production of production of production and proteins of production of productions in a patient's genome that affect the patients own content of contains of proteins or content of proteins or content of proteins or content of proteins or content of the properties of polypeptides. For content of the proteins or content of the proteins or content of the proteins or content of proteins or content of polypeptides. For content of polypeptides or content of polypeptides or content of polypeptides or content of polypeptides. For content of polypeptides or content of polypeptides or content of polypeptides or content of polypeptides. For content of polypeptides or content of polypeptides or content of polypeptides or content of polypeptides. Additionally, (I) and its complementary content of polypeptides of similar nucleic acids in samples, and content of polypeptides of similar nucleic acids in samples, and content of polypeptides of continuous proteins of polypeptides. The antibodies may also be used to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic content of polypeptides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                     Human; single nucleotide polymorphism; SNP; genome; gene ther protein therapy; vaccine; probe; diagnostic assay; detection; quantitation; restorative therapy; polymorphic; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-NOV-1999; 99US-0168138P.
29-NOV-2000; 2000US-00726173.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 1; Page 1876; 2653pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Polymorphic nucleic acid sequences, useful in genetic testing and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-356160/37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Shimkets RA, Leach M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    30-NOV-2000; 2000WO-US032758
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     quantitation; restorative therapy; polymorphic;
                                                      Homo sapiens
                                                                                                                                                                Human silent SNP containing nucleic acid SEQ:5329.
                                                                                                                                                                                                     09-NOV-2001
                                                                                                                                                                                                                                                                               AAI78388 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              polypeptides in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (CURA-) CURAGEN CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                           175
                                                                                                                                                                                                                                                                                                                                                                          51
                                                                                                                                                                                                                                                                                                                                                                                                                                                 47;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       51 BP; 18 A; 14 C; 10 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                           TTTTAGTAGAGATGGAGTTTCTCCATGTTGGTCAGGCTGGTCTCGAACTCC 225
                                                                                                                                                                                                                                                                                                                                                                          TTTTAGTAGACATGGGGTTTCACCATGTTGGTCAGGCTGGTCTTGAACTCC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                samples
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4.5%;
                                                                                                                                                                                                                                                                                 51
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 44.6; DB 1; Length 51; Pred. No. 1.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
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WO200140521-A2

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RESULT 47
AAI79819
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 Query Match
Best Local Similarity
Matches 47; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              by expressing inactive proteins or to supplement the patients own production of polypeptide. Additionally, (I) and its complementary sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and therefore which patients may be in need of restorative therapy. The polypeptides encoded by (I) may be used as antigens in the production of antibodies specific for polymorphic polypeptides. The antibodies may also be used to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sequences (I), which contain single nucleotide polymorphisms (SNPs).

AAM53114 to AAM5329 represent peptides related to human polymorphic polymucleotide sequences. The sequences can be used in gene and protein therapy, and in vaccine production. (I) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of polymorphic polypeptides. For example, (I) may be used to treat disorders by rectifying mutations or deletions in a patient's genome that affect the activity of polypeptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 51 BP; 10 A; 19 C; 13 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide sequences (I), which contain single nucleotide polymorphisms (SNPs).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; Page 2141; 2653pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Polymorphic nucleic acid sequences, useful in genetic testing and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-356160/37
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30-NOV-1999; 99US-0168138P
29-NOV-2000; 2000US-00726173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30-NOV-2000; 2000WO-US032758.
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                                                                                                                                                                                                    Human; single nucleotide polymorphism; SNP; genome; gene therapy; protein therapy; vaccine; probe; diagnostic assay; detection; quantitation; restorative therapy; polymorphic; ds.
                                                                                                                                                                                                                                                                             Human nonconservative amino acid changing SNP nucleic acid SEQ:6760
                                                                                                                                                                                                                                                                                                                                                                                               AAI79819 standard; DNA; 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  polypeptides in samples
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (CURA-) CURAGEN CORP.
30-NOV-1999; 99US-0168138P.
29-NOV-2000; 2000US-00726173.
                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                                                      09-NOV-2001 (first entry)
                                                         30-NOV-2000; 2000WO-US032758
                                                                                              07-JUN-2001
                                                                                                                                   WO200140521-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   847 CCTCGGCCTCCCAAAGTGCTGGGATTACAGGCGTGAGCCACCACGCCCGGC 897
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 CCTCGGCCTCCCAAAGTGCTAGCATTACAGGCGTGAGCCACCATGCCTGGC 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Leach M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4.5%;
92.2%;
                                                                                                                                                                                                                                                                                                                                                                                                 ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 44.6;
Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            .4e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1; Length 51;
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cc sequences (1), which contain single nucleotide polymorphisms (SNPs).

cc AAM53114 to AAM53329 represent peptides related to human polymorphic cc polynucleotide sequences. The sequences can be used in gene and protein cc therapy, and in vaccine production. (1) and the polymorphic polymorphisms which is a patient's genemate by rectifying mutations or cc example, (1) may be used to treat disorders by rectifying mutations or cdeletions in a patient's genome that affect the activity of polypeptides. For cc by expressing inactive proteins or to supplement the patients own cc production of polypeptide. Additionally, (1) and its complementary cc sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and ct polypeptides encoded by (1) may be used as antigens in the production of polypeptides of similar nucleic acids in samples, and cc polypeptides encoded by (1) may be used as antigens in the production of antibodies specific for polymorphic polypeptides. The antibodies may also be used as antigens in the production of conventions and capital agents for detecting the presence of polymorphic polypeptides.
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                                                                                           Query Match
Best Local (
                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 1; Page 2573; 2653pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Shimkets RA, Leach M;
                                                                                                                                                   Sequence 51 BP; 8 A; 16 C; 15 G; 12 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Polymorphic nucleic acid sequences, useful in genetic testing and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2001-356160/37.
                                                                                                                                                                                            polypeptides in samples
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (CURA-) CURAGEN CORP
                    695 CGGGTTCAAGTTATTCTCCTGCCCCAGCCTCCTGAGTAGCTGGGACTACAG 745
                                                                             47;
  ب
                                                                                               Similarity
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AAI79867
                                                                             Conservative
                                                                                           4.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     represent isolated human polymorphic polynucleotide
                                                                             0
                                                                                               Score 44.6;
Pred. No. 1.
                                                                               Mismatches
                                                                                                 1.4e+02;
                                                                                                                     DB 1;
                                                                               4.
                                                                                                                   Length 51;
                                                                               Indels
                                                                               0;
                                                                               Gaps
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RESULT 48
AAI77677/c
ID AAI77
Human; single nucleotide polymorphism; SNP; genome; gene therapy; protein therapy; vaccine; probe; diagnostic assay; detection; quantitation; restorative therapy; polymorphic; ds.
                                                                                                                                                                                                           Human silent SNP containing nucleic acid SEQ:4618.
                                                                                                                                                                                                                                  09-NOV-2001
                                                                                                                                                                                                                                                      AAI77677;
                                                                                                                                                                                                                                                                          677/c
AAI77677 standard;
                                                           30-NOV-1999; 99US-0168138P
29-NOV-2000; 2000US-00726173
                                                                                        30-NOV-2000; 2000WO-US032758
                                                                                                            07-JUN-2001.
WPI; 2001-356160/37
                    Shimkets RA,
                                                                                                                                                  Homo sapiens.
                                       (CURA-) CURAGEN CORP.
                                                                                                                               WO200140521-A2
                                                                                                                                                                                                                                  (first entry)
                     Leach M;
                                                                                                                                                                                                                                                                          DNA;
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RESULT 49
AAH89407
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       by expressing inactive proteins or to supplement the patients own production of polypeptide. Additionally, (I) and its complementary sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and therefore which patients may be in need of restorative therapy. The polypeptides encoded by (I) may be used as antigens in the production of antibodies specific for polymorphic polypeptides. The antibodies may also be used to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic polypeptides in samples
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide sequences (I), which contain single nucleotide polymorphisms (SNPB). AAM53114 to AAM53129 represent peptides related to human polymorphic polynucleotide sequences. The sequences can be used in gene and protein therapy, and in vaccine production. (I) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of polymorphic polypeptides. For example, (I) may be used to treat disorders by rectifying mutations or deletions in a patient's genome that affect the activity of polypeptides
                                                         Isolated human polynucleotides containing single nucleotide polymorphisms, useful for the treatment and diagnosis of e.g. cancer.
                                                                                                          P-PSDB;
                                                                                                                                                        Shimkets
                                                                                                                                                                                                                    07-JAN-2000; 2000US-0174962P
                                                                                                                                                                                                                                                  05-JAN-2001; 2001WO-US000322.
                                                                                                                                                                                                                                                                                  19-JUL-2001
                                                                                                                                                                                                                                                                                                                 WO200151670-A2
                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                             Human; single nucleotide polymorphism;
                                                                                                                                                                                                                                                                                                                                                                                                                          Human coding sequence polymorphic site
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAH89407 standard; DNA; 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; Page 1924; 2653pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Polymorphic nucleic acid sequences, useful in genetic testing and
                                                                                                                                                                                                                                                                                                                                                                              forensic
                                          infection and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local
                                                                                                                          2001-451871/48
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    51
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                                                                                                          AAM00294.
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                                                                                                                                                    R.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                      Leach MD;
                                             diabetes.
                                                                                                                                                                                                                                                                                                                                                                            aberrant protein
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92.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21 G;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 44.6; DB 1; Length 51; Pred. No. 1.4e+02; O; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                            expression;
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Claim 1;

Page 160; 475pp; English

Sequence

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to human nucleic acids containing single nucleotide polymorphisms (SNPs). These can be used in forensic and paternity tests, and to aid in the treatment of disease associated with aberrant protein expression, including cancer, amyloidosis, diabetes, altheimer's disease, Down's syndrome, oedema, thrombocytopaenia, arthritis, glomerulonephritis, haemolytic anaemia, thrombocytopaenia, arthritis, menningitis, muscular disorders, dementia, neurological diseases, tuberous
                                              The present invention relates to human nucleic acids containing single nucleotide polymorphisms (SNPs). These can be used in forensic and paternity tests, and to aid in the treatment of diseases associated with aberrant protein expression, including cancer, amyloidosis, diabettes, Alzheimer's disease, Down's syndrome, oedema, lupus (SLE), vasculitis, glomerulonephritis, haemolytic anaemia, thrombocytopaenia, arthritis, meningitis, muscular disorders, dementia, neurological diseases, tuberous sclerosis, male infertility, hypercalcaemia, blood pressure disorders, osteoporosis, pathogenic infections, hypercholesterolaemia, obesity or autoimmunity. The present sequence is a polymorphism-containing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           meningitis, muscular disorders, dementia, neurological diseases, tuberous sclerosis, male infertility, hypercalcaemia, blood pressure disorders, osteoporosis, pathogenic infections, hypercholesterolaemia, obesity or autoimmunity. The present sequence is a polymorphism-containing
                                                                                                                                                                                                                                                               Isolated human polynucleotides containing single nucleotide polymorphisms, useful for the treatment and diagnosis of e. infection and diabetes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human coding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAH89406 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               oligonucleotide fragment of the invention
                                  oligonucleotide
                                                                                                                                                                                                                               Claim 1; Page 159; 475pp; English.
                                                                                                                                                                                                                                                                                                                                                                                  Shimkets RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                               07-JAN-2000; 2000US-0174962P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 05-JAN-2001; 2001WO-US000322.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200151670-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; single nucleotide polymorphism; SNP;
forensic test; aberrant protein expression;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAH89406;
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                                                                                                                                                                                                                                                                                                                                                                                                                  (CURA-) CURAGEN CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sapiens.
                                                                                                                                                                                                                                                                                                                                                   2001-451871/48.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 974 CTCACTGCAACCTCTGCCTCCCGGGCTCAAGCGATTCTCCTGTCTCAGCCT 1024
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 Similarity 47; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  <u>ب</u>
                                                                                                                                                                                                                                                                                                                                  AAM00293.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP; 8 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sequence
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                                                                                                                                                                                                                                                                                                                                                                                  Leach
                                  fragment of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA;
                                                                                                                                                                                                                                                                                                                                                                                   ₹
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              23 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               92.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 polymorphic site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  51
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                                   the
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 44.6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12 T; 0 U; 0 Other;
                                   invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               .4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 paternity test; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 51;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                  of e.g. cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
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DB 1;

Length 51;

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RESULT 51
ABL00260
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밁
                            S
                                                                                                                                                   CC comprising one or more single nucleotide polymorphisms (SNPS). ABB56531
CC to ABB56903 represent human peptides encoded by some of the SNP
CC oligonucleotides. The sequences from the present invention can have
immunosuppressive, cytostatic, antiinflammatory, neuroprotective and
cc antimicrobial activities. Nucleic acids, polypeptides, oligonucleotides
cc and antibodies from the present invention can be used for treating a
cc subject suffering from, at risk for, or suspected of, suffering from a
cc pathology ascribed to the presence of a sequence polymorphism. The
cc pathology may be autoimmune diseases, inflammation, cancer, diseases of
cc the nervous system, and infection by pathogenic microorganisms. The
cc polymorphism are present in individuals. The antibodies may be used in
cc polymorphism are present in individuals. The antibodies may be used in
cc polymorphic protein (e.g., for use in measuring levels of the polymorphic
cc protein within appropriate physiological samples)
                                                           Query Match
Best Local S
Matches 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; single nucleotide polymorphism; SNP; polymorphism; cytostatic; immunosuppressive; antiinflammatory; neuroprotective; antimicrobial; autoimmune disease; inflammation; cancer; nervous system disease; infection; polymorphic protein; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABL00260 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Isolated human nucleic acids comprising one or more single nucleotide polymorphisms, useful for treating a subject suffering from a pathology, e.g. autoimmune diseases, ascribed to the presence of a sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     31-MAY-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200138586-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human silent noncoding SNP oligonucleotide SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      05-MAR-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24-NOV-1999; 99US-0167383P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     22-NOV-2000; 2000WO-US032311.
                                                                                                                                                                                                                                                                                                                                                                                                         ABL00010 to ABL01104 represent human nucleic acid oligonucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 1; Page 323; 674pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      polymorphism
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                              954
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                                                             l Similarity
47; Conserv
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                                                                                                                             51
                GTGCAATGGCCAAATCTCGGCTCACTGCAACCTCTGCCTCCCGGGCTCAAG 1004
 GTGCAATGGCATGATCTCGGCTCACTGCAACCTCTGCCTCCCGGGTTCAAG 51
                                                                                                                             BP;
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                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
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                                                                                                                             Α;
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92.2%;
                                                                              4.5%;
92.2%;
                                                                                                                             17 C; 13 G; 12 T; 0 U; 0 Other;
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                                                              Score 44.6; DI
Pred. No. 1.4e.
0; Mismatches
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Pred. No. 1.4e+02;
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                                                                              4e+02
                                                                                              DB 1;
                                                                                             Length 51
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                                                                  Indels
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                                                              0,
                                                                                                                                                                            polymorphic
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                                                                  Gaps
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RESULT 52
AAH89761
ID AAT
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AAH89763
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAH89761;
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                                                                                                                                                                                                                                                                                                 Sequence 50 BP; 10 A; 10 C; 11 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; single nucleotide polymorphism;
forensic test; aberrant protein express
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human coding sequence polymorphic site
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to human nucleic acids containing single nucleotide polymorphisms (SNPs). These can be used in forensic and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1; Page 260; 475pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Isolated human polynucleotides containing single nucleotide polymorphisms, useful for the treatment and diagnosis of e.g. cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Shimkets
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               07-JAN-2000; 2000US-0174962P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
                                                                                                AAH89763
                                                                                                                        AAH89763 standard; DNA; 50 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          infection and diabetes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (CURA-) CURAGEN CORP.
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                                                                                                                                                                                                                    750 CCACCACGCCTAGCTAATTTTTTGTATTTTTAGTAGAGATGGGGTT
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                                                                                                                                                                                                                                                Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Leach
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3
                                                                                                                                                                                                                                                               4.4%;
95.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   e polymorphism; SNP;
protein expression;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 50
                                                                                                                                                                                                                                                0,
                                                                                                                                                                                                                                                               Score 43.8; DB 1;
Pred. No. 1.5e+02;
                                                                                                                                                                                                                                                  Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       paternity test;
                                                                                                                                                                                                                                                                            Length
                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                         tuberous
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Human; single nucleotide polymorphism; SNP; forensic test; aberrant protein expression;

paternity test; ds.

Human coding 01-OCT-2001

(first entry)

0

sequence polymorphic site SEQ ID NO: 544

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ID XXX ACC XXX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity
Matches 45; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    nucleotide polymorphisms (SNPs). These can be used in forensic and paternity tests, and to aid in the treatment of diseases associated with aberrant protein expression, including cancer, amyloidosis, diabetes, Alzheimer's disease, Down's syndrome, oedema, lupus (SLE), vasculitis, glomerulonephritis, haemolytic anaemia, thrombocytopaenia, arthritis, meningitis, muscular disorders, dementia, neurological diseases, tuberous sclerosis, male infertility, hypercalcaemia, blood pressure disorders, osteoporosis, pathogenic infections, hypercholesterolaemia, obesity or autoimmunity. The present sequence is a polymorphism-containing oligonucleotide fragment of the invention
Shimkets RA,
                                                                                                 07-JAN-2000;
                                                                                                                                                       05-JAN-2001;
                                                                                                                                                                                                        19-JUL-2001.
                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                        Human; single nucleotide polymorphism; SNP;
                                                                                                                                                                                                                                                                                                                                                                                                                                            Human coding sequence polymorphic site
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAH89759 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1; Page 260; 475pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             polymorphisms, userur rinfection and diabetes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Isolated human polynucleotides containing single nucleotide polymorphisms, useful for the treatment and diagnosis of e.g. cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Shimkets RA,
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                                                                                                                                                                                                                                                         WO200151670-A2
                                                                                                                                                                                                                                                                                                                                                                   forensic
                                                   (CURA-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (CURA-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           present invention relates to human nucleic acids containing single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2001-451871/48.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      750
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      N
                                                   CURAGEN CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAM00646.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CCACCACGCCTAGCTAATTTTTTTGTATTTTTAGTAGAGATGGGGTT 796
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         50 BP; 10
                                                                                                                                                                                                                                                                                                                                                                test;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                    2000US-0174962P
                                                                                                                                                       2001WO-US000322
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Leach MD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
Leach MD;
                                                                                                                                                                                                                                                                                                                                                                aberrant protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; 10 C; 11 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4.4%;
95.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       <u>.</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score
Pred.
                                                                                                                                                                                                                                                                                                                                                                expression;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                43.8;
No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            рь
1.5e+02;
2;
                                                                                                                                                                                                                                                                                                                                                                   paternity test, ds.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 540
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RESULT 55
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Matches 45; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                aberrant protein expression, including cancer, amyloidosis, diabetes, Alzheimer's disease, Down's syndrome, oedema, lupus (SLE), vasculitis, glomerulonephritis, haemolytic anaemia, thrombocytopaenia, arthritis, meningitis, muscular disorders, dementia, neurological diseases, tuberous sclerosis, male infertility, hypercalcaemia, blood pressure disorders, osteoporosis, pathogenic infections, hypercholesterolaemia, obesity or autoimmunity. The present sequence is a polymorphism-containing oligonucleotide fragment of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention relates to human nucleic acids containing single nucleotide polymorphisms (SNPs). These can be used in forensic and paternity tests, and to aid in the treatment of diseases associated with
AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide sequences (I), which contain single nucleotide polymorphisms (SNPs).

AAM53114 to AAM53329 represent peptides related to human polymorphic polynucleotide sequences. The sequences can be used in gene and protein
                                                                                                                                                                                                                                                                                                                                                                                                             Human; single nucleotide polymorphism; SNP; genome; protein therapy; vaccine; probe; diagnostic assay; quantitation; restorative therapy; polymorphic; ds.
                                                                                                                                                                                                                                            30-NOV-1999;
29-NOV-2000;
                                                                                                                                                                                                                                                                                           30-NOV-2000; 2000WO-US032758
                                                                                                                                                                                                                                                                                                                         07-JUN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           09-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   polymorphisms, useful f
infection and diabetes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Isolated human polynucleotides containing single nucleotide polymorphisms, useful for the treatment and diagnosis of e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                P-PSDB; AAM00642.
                                                                         Claim 1;
                                                                                                                     Polymorphic nucleic acid
                                                                                                                                                   WPI; 2001-356160/37.
                                                                                                                                                                                                                                                                                                                                                       WO200140521-A2
                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human silent SNP containing nucleic acid SEQ:2541.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAI75600
                                                                                                                                                                                                                (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2001-451871/48.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               750 CCACCACGCCTAGCTAATTTTTTTGTATTTTAGTAGAGATGGGGTT 796
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                                                                       Page
                                                                                                                                                                                  ŖΑ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CCACCACGCCTGGCTAATTTTTTGTATTTTTAGTAGAGACGGGGTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      50 BP; 10 A; 11 C; 11 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      standard; DNA; 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                              99US-0168138P.
2000US-00726173.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                         829; 2653pp;
                                                                                                                                                                                  Leach
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4.4%;
95.7%;
                                                                                                                     sequences,
                                                                         English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 43.8;
Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                       useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          .5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1;
                                                                                                                     in genetic testing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                  detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                              gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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RESULT 56
AD11250/c
ID AD1125
XX AD1125
XX AD1125
XX AD125
XX AD12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ś
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local S
Matches 45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  therapy, and in vaccine production. (I) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of polymorphic polypeptides. For example, (I) may be used to treat disorders by rectifying mutations or deletions in a patient's genome that affect the activity of polypeptides by expressing inactive proteins or to supplement the patients own production of polypeptide. Additionally, (I) and its complementary sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and therefore which patients may be in need of restorative therapy. The polypeptides encoded by (I) may be used as antigens in the production of antibodies specific for polymorphic polypeptides. The antibodies may also be used to down regulate expression and activity. The antibodies may also be used to down regulate expression and activity. The antibodies may also be used disamnetic access for activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADI12550
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADI12550 standard;
This invention relates to a novel method for predicting a predisposition to cancer in a patient by detecting large deletions in the human tumour suppressor gene identified as the breast cancer susceptibility gene 1 (BRCA1). Specifically, it refers to deletions that result from the unequal crossover between a pair of repetitive Alu sequences in the BRCA1 gene, such that the recombined nucleotide sequence containing the deletion indicates a predisposition to breast and ovarian cancer. The present invention describes newly discovered deletion mutations that are believed to be deleterious and cause significant alterations in the
                                                                                                                                                                                                                                                                                                                                                        predicting a predisposition to cancer in a patient comprising deletion in the BRCA1 gene that results from the unequal cross between a pair of repetitive sequences in the BRCA1 gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mutant human BRCA1 genomic DNA resulting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  07-JUN-2002;
09-AUG-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ovarian cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ds; cancer; human; tumour suppressor;
breast cancer susceptibility gene 1; BRCA1; repetitive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        22-APR-2004
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                                                                                                                                                                                                                                                                                                      Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2004-062369/06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Scholl T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             09-JUN-2003; 2003WO-US018098
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO2003104474-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           634
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MYRIAD GENETICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ACTCTGTCACCCAGGCTGGAGTGCAGTGGCGCAATCTTGGCTCACTG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hendrickson BC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP; 13
                                                                                                                                                                                                                                                                                                      SEQ ID NO 33;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2002US-0387132P
2002US-0402430P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           recombination; mutant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4.4%;
95.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               50
                                                                                                                                                                                                                                                                                                   59pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ward
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score
Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                      English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              В
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  43.8;
No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pruss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 1;
.5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  from deletion 5 SeqID
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
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                                                                                                                                                                                                                                                                                                                                                                                                 crossover
                                                                                                                                                                                                                                                                                                                                                                                                                                detecting
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                                                                          structure or biochemical function of BRCA1. Accordingly, it provides methods for detecting such mutants, as well as identifying and screening for cytostatic compounds useful for treating or preventing cancers associated with a BRCA1 genetic variant. This polynucleotide is a mutant human BRCA1 genomic DNA fragment that arises as a result of a recombination event (deletion 5), which causes the omission of exons 15
                                                     given in an exemplification of the invention.
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Sequence
               50 BP; 13
               A; 12 C;
                15
                G; 10 T; 0 U; 0 Other;
BB
Length
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Matches
                                 Query Match
Best Local Similarity
     695
                        46;
CGGGTTCAAGTTATTCTCCTGCCCCAGCCTCCTGAGTAGCTGGGACTACA
                         Conservative
                                 4.4%;
                          <u>.</u>
                                 Score 43.6;
Pred. No. 1
                          Mismatches
                                    1.6e+02;
                          4.
                           Indels
                          0
                          Gaps
                            0
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AAL29844 standard; DNA; 51 뫄 밁 S

50

CGGGTTCAAGCAATTCTCCTGCCTCAGCCTCCTGAGTAGCTGGGATTACA

AAL29844;

24-JAN-2002 (first entry)

Human SNP oligonucleotide #3052

Immunosuppressive; immunostimulatory; antiinflammatory; cytostatic; neuroprotective; antimicrobial; gene therapy; vaccine; amylase; cancer amyloid protein; angiopoietin; apoptosis related protein; cadherin; cyclin; polymerase; oncogene; histone; kinase; colony stimulating factory complement related protein; cytochrome; kinase; cytokine; interferon; complement related protein; cytochrome; kinesin; cytokine; interferc interleukin; G-protein coupled receptor; thioesterase; inflammation; multifactorial disease; autoimmune disease; infection; nervous system disease; ss. system disease; cancer; factor;

iomo sapiens.

WO200147944-A2

28-DEC-2000; 2000WO-US035498

28-DEC-1999; 99US-0173419P. 27-DEC-2000; 2000US-00173419.

(CURA-) CURAGEN CORP.

Shimkets RA, WPI; 2001-465210/50 Leach 3

Polymorphic nucleic acids encoding e.g. amylases, cyclins, polymerases, oncogenes and histones, useful for diagnosing and treating, e.g. cancer autoimmune diseases and infections.

Claim 1; Page 2260; 4143pp; English.

RESULT 57
RAL29844
ID PAL228
XX AL22
XX AL22
XX AL22
XX Inmun
XX I The present invention relates to oligonucleotides encoding polymorphic variants of proteins related to amylases, amyloid proteins, angiopoietir apoptosis related proteins, candherin, cyclin, polymerase, oncogenes, histones, kinases, colony stimulating factors, complement related proteins, cytochromes, kinesins, cytokines, interferons, interleukins, opportein coupled receptors and thioesterases. The present sequence is one such oligonucleotide. The oligonucleotides and the peptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of the proteins listed above. Disorders that may be prevented, diagnosed and/or treated include multifactorial diseases with a genetic component, such as autoimmune diseases (e.g. rheumatoid diabetes angiopoietin one Ģ

Length 51;

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RESULT 58
AAI73532/
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                                          cc sequences (I), which contain single nucleotide polymorphisms (SNPs).

C AAM53114 to AAM5329 represent peptides related to human polymorphic polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases considered with inappropriate expression of polymorphic polypeptides. For conditions in a patient's genome that affect the activity of polypeptides or conditions in a patient's genome that affect the activity of polypeptides by expressing inactive proteins or to supplement the patients own considered production of polypeptide. Additionally, (I) and its complementary considered patients may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and considered patients may be used as antigens in the production of polypeptides encoded by (I) may be used as antigens in the production of considered patients of polymorphic polypeptides. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic conditions in samples and activity. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic conditions and activity.
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Matches
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29-NOV-2000; 2000US-00726173.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   systemic lupus erythromatosus and Grave's disease), inflammation, cancer (e.g. cancers of the bladder, brain, breast, colon and kidney, leukaemia), diseases of the nervous system and an infection of pathogenia
                                                                                                                                                                                                                                                                                                                                                                         AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Polymorphic nucleic acid sequences, useful in genetic testing and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (CURA-) CURAGEN CORP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Leach M;
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Pred. No. 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQ: 473
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51

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> Matches Query Match Best Local (

46; Similarity

Conservative

0

4.4%;

Score 43.6; D Pred. No. 1.6e 0; Mismatches

6e+02

Length 51

Indels

Gaps

0

695 0;

646 AGGCTGGAGTGCAGTGGCGCAATCTTGGCTCACTGCAACCTCTGCCTCCC

Sequence

51 BP; 11 A; 14 C; 17 G; 9 T; 0 U; 0 Other;

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RESULT 59
AAI79585/c
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Best Local Similarity
                                                                                                      polymucleotide sequences. The sequences can be used in gene and protein therapy, and in vaccine production. (I) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of polymorphic polypeptides. For example, (I) may be used to treat disorders by rectifying mutations or deletions in a patient's genome that affect the activity of polypeptides by expressing inactive proteins or to supplement the patients own production of polypeptide. Additionally, (I) and its complementary sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and therefore which patients may be in need of restorative therapy. The polypeptides encoded by (I) may be used as antigens in the production of
                                                   antibodies specific for polymorphic polypeptides. The antibodies may also be used to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic
                                                                                                                                                                                                                                                                                                                                    AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide sequences (I), which contain single nucleotide polymorphisms (SNPs). AAM53114 to AAM53329 represent peptides related to human polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; single nucleotide polymorphism; SNP; genome; gene ther protein therapy; vaccine; probe; diagnostic assay; detection; quantitation; restorative therapy; polymorphic; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                Claim 1; Page 2504; 2653pp; English
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29-NOV-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   09-NOV-2001
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                                 polypeptides in samples
                                                 be used as diagnostic agents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Polymorphic nucleic acid sequences, useful in genetic testing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-356160/37
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2000US-00726173.
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Pred. No. 1.6e+02;
0; Mismatches 4
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RESULT 60
RABIT455/c
ID AAIT455/c
ID AAIT455/c
ID AAIT455/c
ID AAIT455/c
ID AAIT455/c
ID AAIT455/c
ID O9-NOV
XX Human
XX Human
XX Human
XX Human
XX Quanti
XX Quanti
XX Quanti
XX Guanti
XX Guanti
XX Guanti
XX Guanti
XX Guanti
XX Guanti
XX JO-NOV
XX JO-NOV
XX JO-NOV
XX JO-NOV
XX GURAXX Human
CC CURAXX Guanti
CC AAIT31
CC AAIT31
CC AAIT31
CC DAIM
CC Lherag
CC AAIT31
CC DAIM
CC AAIT31
CC DAIM
CC C AAIT31
CC DAIM
CC AAIT31
CC DAIM
CC C AAIT31
CC DAIM
CC C AAIT31
CC DAIM
CC C AAIT31
CC DAIM
CC D AAI74555 standard; DNA; 51 AAI74555; 09-NOV-2001 (first entry) ₽P

Human silent SNP containing nucleic acid SEQ:1496.

Human; single nucleotide polymorphism; SNP; genome; protein therapy; vaccine; probe; diagnostic assay; quantitation; restorative therapy; polymorphic; ds. ; gene therapy; detection;

07-JUN-2001 WO200140521-A2

Homo sapiens

30-NOV-2000; 2000WO-US032758

30-NOV-1999; 99US-0168138P. 29-NOV-2000; 2000US-00726173.

(CURA-) CURAGEN CORP.

Shimkets RA, WPI; 2001-356160/37 Leach M;

Polymorphic nucleic acid sequences, useful in genetic testing and

Claim 1; Page 511; 2653pp; English

by expressing inactive proteins or to supplement the patients own production of polypeptide. Additionally, (I) and its complementary sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and therefore which patients may be in need of restorative therapy. The polypeptides encoded by (I) may be used as antigens in the production of antibodies specific for polymorphic polypeptides. The antibodies may also be used to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic polypeptides in samples AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide sequences (I), which contain single nucleotide polymorphisms (SNPB). AAM53114 to AAM53229 represent peptides related to human polymorphic polynucleotide sequences. The sequences can be used in gene and protein therapy, and in vaccine production. (I) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of polymorphic polypeptides. For example, (I) may be used to treat disorders by rectifying mutations or deletions in a patient's genome that affect the activity of polypeptides.

Sequence 51 BP; 10 A; 14 C; 17 G; 10 T; 0 U; 0 Other;

밁 Ś Matches Query Match Best Local 635 50 Similarity CTCTGTCACCCAGGCTGGAGTGCAGTGGCGCAATCTTGGCTCACTGCAAC 684 CTCTGTCACCCAGGCTGGAGTGCAATGGCACGATCTCGGCTCACTGCAAC 1 4.4%; larity 92.0%; Conservative Score 43.6; DB 1; Length 51; Pred. No. 1.6e+02; O; Mismatches 4; Indels 0 Gaps

0

밁 Ś

638 TGTCACCCAGGCTGGAGTGCAGTGGCGCAATCTTGGCTCACTGCAACCTC 687 TGTCACCCAGGCTGAACTGCAGTGGCGTGATCTTGGCTCACTGCAACCTC

50

Gaps

0

RESULT 62
AAI77807
ID AAI77
XX
AC AAI77
XX
AC AAI77
XX
DT 09-NC

AAI77807 standard; DNA; 51 BP

09-NOV-2001 AAI77807;

(first entry)

RESULT 61 AAI73861

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Matches
                                                     Query Match
Best Local :
                                                                                                                                                                                                                                       associated with inappropriate expression of polymorphic polypeptides. For example, (I) may be used to treat disorders by rectifying mutations or deletions in a patient s genome that affect the activity of polypeptides by expressing inactive proteins or to supplement the patients own production of polypeptide. Additionally, (I) and its complementary sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and therefore which patients may be in need of restorative therapy. The polypeptides encoded by (I) may be used as antigens in the production of antibodies specific for polymorphic polypeptides. The antibodies may also be used to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic polymorphicals.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  polynucleotide sequences. The sequences can be used in gene and protein therapy, and in vaccine production. (I) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases
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29-NOV-2000; 2000US-00726173.
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                                                                                                                                                     Sequence 51 BP; 9 A; 17 C; 13 G; 12 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1; Page 299; 2653pp; English.
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                                                                                                                                                                                                                      polypeptides in samples
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Local Similarity 92.0%; hes 46; Conservative
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                     0
                                                     Score 43.6; DB 1;
Pred. No. 1.6e+02;
                        Mismatches
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                                                                                            DB 1;
                                                                                            Length 51;
                            Indels
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Human silent SNP

containing nucleic acid SEQ:4748

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AAI77520/c
ID AAI775
XX AAI775
AC AAI775
XX D9-NOV
DT 09-NOV
XX Human
XX Human;
KW Human;
KW granti
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     CC AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide CC sequences (I), which contain single nucleotide polymorphisms (SNPs). CC AAM53114 to AAA53329 represent peptides related to human polymorphic CC polynucleotide sequences. The sequences can be used in gene and protein CC therapy, and in vaccine production. (I) and the polypeptides encoded by C them may be used in the prevention, diagnosis and treatment of diseases CC associated with inappropriate expression of polymorphic polypeptides. For CC deletions in a patient's genome that affect the activity of polypeptides or CC deletions in a patient's genome that affect the activity of polypeptides by expressing inactive proteins or to supplement the patients own CC production of polypeptide. Additionally, (I) and its complementary CC sequences may also be used as DNA probbes in diagnostic assays to detect CC and quantitate the presence of similar nucleic acids in samples, and CC therefore which patients may be in need of restorative therapy. The CC polypeptides encoded by (I) may be used as antigens in the production of CC polypeptides encoded by (I) may be used as antibodies may also be used to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic
                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
Matches 46; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo
                 Human; single nucleotide polymorphism; SNP; genome; gene therapy; protein therapy; vaccine; probe; diagnostic assay; detection; quantitation; restorative therapy; polymorphic; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2001-356160/37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Shimkets RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      30-NOV-1999; 99US-0168138P.
29-NOV-2000; 2000US-00726173.
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                                                                                                                                 09-NOV-2001
                                                                                                                                                                                                         AAI77520 standard; DNA; 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; Page 1963; 2653pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      polypeptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Polymorphic nucleic acid sequences, useful in genetic testing and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (CURA-)
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                                                                                                                                                                                                                                                                                                                                         672 GGCTCACTGCAACCTCTGCCTCCCGGGTTCAAGTTATTCTCCTGCCCCAG 721
                                                                                            silent
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                                                                                                                                                                                                                                                                                                    GGCTCACTGCATCCTCCGCCTCCCGGGTTCAAGCTATTCTCCTGCCTCAG 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 6 A; 22 C; 10 G; 13 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Leach M;
                                                                                            SNP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      in samples
                                                                                                                             (first entry)
                                                                                         containing nucleic
                                                                                                                                                                                                                                                                                                                                                                                           4.4%;
                                                                                                                                                                                                         BP
                                                                                                                                                                                                                                                                                                                                                                            0,
                                                                                                                                                                                                                                                                                                                                                                                           Score 43.6; DB 1;
Pred. No. 1.6e+02;
                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                            acid
                                                                                            SEQ:4461.
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                                                                                                                                                                                                                                                                                                                                                                                                             Length 51;
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                                                                                                                                                                                                                                                                                                                                                                          0;
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CC AAM53114 to AAM53329 represent peptides related to human polymorphic cyclophynucleotide sequences. The sequences can be used in gene and protein cyclophynucleotide sequences. The sequences can be used in gene and protein cyclophynucleotide sequences. The sequences can be used in gene and protein cyclophynucleotide sequences can be used in gene and protein cyclophynucleotides encoded by them may be used in the prevention, diagnosis and treatment of diseases cascolated with inappropriate expression of polymorphic polymorphic polypeptides. For cyclophynucleotides in a patient's genome that affect the activity of polypeptides or cyclophynucleotides over cyclophynucleotides in samples, and cyclophynucleotides encoded by (I) may be used as antigens in the production of cyclophynucleotides encoded by (I) may be used as antigens in the production of cyclophynucleotides encoded by (I) may be used as antigens in the production of cyclophynucleotides encoded by (I) may be used as antigens in the production of cyclophynucleotides encoded by (I) may be used as antigens in the production of cyclophynucleotides encoded by (I) may be used as antigens in the production of cyclophynucleotides encoded by (I) may be used as antigens in the production of cyclophynucleotides encoded by (I) may be used as cyclophynucleotides encoded by (I) may be used as antigens in the production of cyclophynucleotides encoded by (I) may be used as antigens in the production of cyclophynucleotides encoded by (I) may be used as antigens in the production of cyclophynucleotides encoded by (I) may be used as cyclophynucleotides encode
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Shimkets RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30-NOV-1999; 99US-0168138P
29-NOV-2000; 2000US-00726173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-NOV-2000; 2000WO-US032758
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    polypeptides in samples
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Polymorphic nucleic acid sequences, useful in genetic testing
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    46;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                                                 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TTTAGTAGACATGGGGTTTCACCATGTTGGTCAGGCTGGTCTTGAACTCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TTTAGTAGAGATGGAGTTTCTCCATGTTGGTCAGGCTGGTCTCGAACTCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          51 BP; 18 A; 14 C; 10 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4.4%;
92.0%;
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Pred. No. 1.6e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.6e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 51;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genotyping; agammaglobulinaemia; diabettes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis;

WO200129262-A2 Homo sapiens. inflammation;

forensic

investigation; paternity analysis;

14-AUG-2001

(first entry)

0,

Human SNP flanking oligonucleotide SEQ ID 1304.

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AAI73065/c
ID AAI730
XX
AC AAI730
AC AAI730
DY 09-NOV
XX
OF Human
XX
KW Human;
KW protei
KW quanti
XX
COS Homo 8
XX
PD 07-JUN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CC Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide primer extension (SNPB) primers, and the sequences of regions flanking control of the single nucleotide polymorphisms SNPs. The present invention control of sites of single nucleotide polymorphisms SNPs. The present invention control of the invention. The presence of a SNP, using the cligonucleotides of the invention. The PCR primers are used to amplify a contigonucleotides are useful for genotyping a nucleic acid sample by performing a single-nucleotide primer extension reaction. The contigonucleotides are useful for determining the presence, absence or contigonucleotides are useful for determining the presence, absence or contigonucleotides are useful for determining the presence, absence or contigonucleotides are useful for determining the presence, absence or conditional part of control of the presence of
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                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 65
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Best Local S
Matches 46
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              inflammation, cancer, nervous system diseases and infection by pathogenic microorganism. The method is also useful in forensic investigations and paternity analysis. The present sequence represents a fragment of human DNA flanking the site of a single nucleotide polymorphism
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; Page 56; 83pp; English
                                                                                                                                         Human; single nucleotide polymorphism; SNP; genome; gene therapy; protein therapy; vaccine; probe; diagnostic assay; detection; quantitation; restorative therapy; polymorphic; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
     07-JUN-2001
                                                                                                                                                                                                                                      Human silent SNP containing nucleic acid SEQ:6.
                                                                                                                                                                                                                                                                                        09-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                  AAI73065 standard; DNA; 51 BP
                                               WO200140521-A2
                                                                                            Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP; 9
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                                                                                                                                                                                                                                                                                      (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16 C; 13 G; 13 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 43.6;
Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ..6e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
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S
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Cc AAM33114 to AAM53329 represent peptides related to human polymorphic CC polymucleotide sequences. The sequences can be used in gene and protein CC therapy, and in vaccine production. (I) and the polymorphic cot them may be used in the prevention, diagnosis and treatment of diseases cas associated with inappropriate expression of polymorphic polymorphic cot example, (I) may be used to treat disorders by rectifying mutations or cot deletions in a patient's genome that affect the activity of polypeptides for polymorphic polymor
                                                                                                        Query Match
Best Local S
Matches 45
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29-NOV-2000; 2000US-00726173
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                                                                                                                                                                                                                   Sequence 51 BP; 13 A; 10 C;
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                                                                                                                                                                                                                                                                      polypeptides in samples
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                                             966 AATCTCGGCTCACTGCAACCTCTGCCTCCCGGGCTCAAGCGATTCTCC 1013
  48
                                                                                                                                    Similarity
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ANTOTOGGCTCACTGCAACCTCCGCCTCCTGGGTTCAAGCGATTCTCC 1
                                                                                                           4.4%;
llarity 93.8%;
Conservative
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                                                                                                              Pred.
0; Mis
                                                                                                                                                                                                                         20 G; 8 T;
                                                                                                                                       Score 43.2;
Pred. No. 1.
                                                                                                                Mismatches
                                                                                                                                                                                                                         0 U; 0 Other;
                                                                                                                                       .7e+02;
                                                                                                                                                                         DB
                                                                                                                                                                1; Length 51;
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                                                                                                                   Indels
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                                                                                                                0;
                                                                                                                   Gaps
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RESULT 66
AAI77325
ID AAI77
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                                                                                                                 Human, single nucleotide polymorphism; SNP; genome; gene ther protein therapy; vaccine; probe; diagnostic assay; detection; quantitation; restorative therapy; polymorphic; ds.
                                                                                                                                                           Human silent
                                                                                                                                                                              09-NOV-2001
                                                                                                                                                                                                                     AAI77325 standard;
        30-NOV-1999; 99US-0168138P
29-NOV-2000; 2000US-00726173
                                     30-NOV-2000; 2000WO-US032758
                                                         07-JUN-2001.
                                                                            WO200140521-A2
                                                                                                                                                          SNP containing nucleic acid SEQ: 4266.
                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                       DNA;
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                                                                                                                                                                                                                       ВÞ
                                                                                                                                         gene therapy;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local S
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Shimkets RA,
                                                          17-NOV-1998;
16-NOV-1999;
                                                                                                                                     25-MAY-2000.
                                                                                                                                                                                                              variation
                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                       Human; single nucleotide polymorphism; SNP; chromosome 8; identification; gene therapy; ss.
                                                                                                                                                                                                                                                                                                                                  Human clone cg43972482 polymorphic site, SEQ ID NO:914.
                                                                                                                                                                                                                                                                                                                                                                     16-NOV-2000
                                                                                                                                                                                                                                                                                                                                                                                                   AAA77231;
                                                                                                                                                                                                                                                                                                                                                                                                                               AAA77231 standard; cDNA; 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1;
                              (CURA-) CURAGEN CORP
                                                                                                       17-NOV-1999;
                                                                                                                                                                 WO200029623-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             polypeptides in samples
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Polymorphic nucleic acid sequences, useful in genetic testing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (CURA-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             843 CCTGCCTCGGCCTCCCAAAGTGCTGGGATTACAGGCGTGAGCCACCAC 890
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   45;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 51
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CCCGCCTCGGCCTCCCAAAGTGTTGGGATTACAGGCGTGAGCCACCGC 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Page 1815; 2653pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP; 8 A; 20 C; 14 G; 9 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                 (first entry)
Leach MD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Leach M;
                                                          98US-0109024P.
99US-00443199.
                                                                                                       99WO-US027293
                                                                                                                                                                                            Location/Qualifiers replace(26,T) /*tag= a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 93.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                               ΒP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 43.2; DB 1; Length 51; Pred. No. 1.7e+02; 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0 Other;
                                                                                                                                                                                                                                                                                                        detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ٥,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
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changes, while those in sequences 1129 to 1186 (AAA77446-A77503) result in non- conservative changes. The SNPs in sequences 1187 to 1192 (AAA77504-A77509) generate frameshift mutations. The invention also relates to a method of detecting a polymorphic site in a nucleic acid and a method of determining the relatedness of two nucleic acids. It also encompasses peptides containing polymorphic sites, antibodies raised against such peptides, and a method of detecting polymorphic proteins/peptides using the antibodies. The nucleic acids are useful for gene therapy of an individual having, suspected of having, or at risk of developing a pathological condition due to the presence of a sequence polymorphism. Such treatment would comprise administration of the wild-type nucleic acid sequence. Antibodies raised against polymorphic peptides can also be used in the treatment of such individuals
                                                                                                                                                                                                                                                                                                                                                                                                                               Sequences AAA76318-A77509 represent 1192 human nucleic acid sequences which contain single nucleotide polymorphisms (SNPs). Sequences 1 to 1112 (AAA76318-A77429) are consecutive pairs of nucleotides which contain silent SNPs. Sequences 1113 to 1192 (AAA77430-A77509) are consecutive pairs of nucleotides containing SNPs which result in changes in the corresponding amino acid sequences (AAB11749-B1128). The SNPs in sequences 1113 to 1128 (AAA77430-A77445) lead to conservative amino acid sequences (ABB1749-B1128).
Sequence 51 BP; 9 A; 12 C; 16 G; 14 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 1; Page 433; 543pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        presence of a sequence polymorphism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human nucleic acids containing single nucleotide polymorphisms, useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2000-387826/33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   for treating a subject suffering,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      or at risk from a pathology due to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1112
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밁 á Matches Query Match Best Local 177 TTAGTAGAGATGGAGTTTCTCCATGTTGGTCAGGCTGGTCTCGAACTCCCG 227 Н 1 Similarity 46; Conserv TTAGTAGAGACGGGGTTTCACCATGCTGGTCAGGCTGGTCTCGAACTCCTG 51 Conservative 4.3%; <u>,</u> Score 43; DB 1; Le Pred. No. 1.7e+02; 0; Mismatches 5; Length 51; Indels , , Gaps 0,

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RESULT 68
AAA77499
ID AAA77
WO200029623-A2
                                                                                                                                       Homo sapiens.
                                                                                                                                                     Human; single nucleotide polymorphism; SNP; detection; identification;
gene therapy; ss.
                                                                                                                                                                             Human Alu subfamily SQ gene polymorphic site, SEQ ID NO:1182.
                                                                                                                                                                                             16-NOV-2000 (first entry)
                                                                                                                                                                                                                            AAA77499 standard; cDNA; 51 BP
                                                                           25-MAY-2000
                                                                                                                  variation
                                                                                                                 replace(26,C)
                                                                                                                         Location/Qualifiers
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Shimkets RA,

Leach MD

CURA-) CURAGEN CORP

17-NOV-1998; 16-NOV-1999; 17-NOV-1999;

98US-0109024P. 99US-00443199. 99WO-US027293

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RESULT 69
RAL31284/c
ID AAL312
XX AAL312
XX AAL312
XX IMMUNO
XX IMMUNO
KW IMMUNO
KW IMMUNO
KW IMMUNO
KW CYClin
KW CYClin
KW CYClin
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XX PPN (CURA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequences AAA76318-A77509 represent 1192 human nucleic acid sequences CC which contain single nucleotide polymorphisms (SNPs). Sequences 1 to 1112 (AAA76318-A77429) are consecutive pairs of nucleotides which contain CC silent SNPs. Sequences 113 to 1192 (AAA77430-A77509) are consecutive CC pairs of nucleotides containing SNPs which result in changes in the CC corresponding amino acid sequences (AAA1749)-B11828). The SNPs in CC sequences 1113 to 1128 (AAA77430-A77445) ladd to conservative amino acid changes, while those in sequences 1129 to 1186 (AAA77446-A77503) result CC (AAA77504-A77509) generate frameshift mutations. The invention also CC relates to a method of detecting a polymorphic site in a nucleic acid and CC amethod of detecting a polymorphic site in a nucleic acid and CC against such peptides containing polymorphic sites, antibodies raised CC gene therapy of an individual having, suspected of having, or at risk of CC developing a pathological condition due to the presence of a sequence CC polymorphism. Such treatment would comprise administration of the wild-ctype nucleic acid sequence. Antibodies raised against polymorphic CC type nucleic acid sequence. Antibodies raised against polymorphic CC conditions are useful for CC developing a pathological condition due to the presence of a sequence CC polymorphism. Such treatment would comprise administration of the wild-ctype nucleic acid sequence. Antibodies raised against polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        밁
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Best Local S
Matches 46
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P-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; Page 515; 543pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human nucleic acids containing single nucleotide polymorphisms, useful for treating a subject suffering, or at risk from a pathology due to t presence of a sequence polymorphism.
                                                                                                                                                                                                                                                                                                      interleukin; G-protein coupled receptor; thioesterase; inflammation; multifactorial disease; autoimmune disease; infection; nervous system disease; ss.
                                                                                                                                                                                                                                                                                                                                                                               neuroprotective; antimicrobial; gene therapy; vaccine; amylase; cancer; amyloid protein; angiopoietin; apoptosis related protein; cadherin; cyclin; polymerase; oncogene; histone; kinase; colony stimulating factor; complement related protein; cytochrome; kinesin; cytokine; interferon;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAL31284 standard; DNA; 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 51
                                                                                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Immunosuppressive; immunostimulatory; antiinflammatory; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human SNP oligonucleotide #4492.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   24-JAN-2002
                                                                   28-DEC-1999; 99US-0173419P.
27-DEC-2000; 2000US-00173419.
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                                                                                                                                       28-DEC-2000; 2000WO-US035498
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)B; AAB11818.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                355 CTGAGCTCAAGCAGTCCACCTGCCTCAGCCTCCCAAAGTGCTGGGATTACA 405
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ب
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               46;
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                         CURAGEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CTGACCTCAAGTGATCCACCTGCCTTAGCCTCCCAAAGTGCTGGGATTACA 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP; 12 A; 17 C; 10 G; 12 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ₽P
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Pred. No. 1.7e+02;
0; Mismatches 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 51;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
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밁 S

Query Match Best Local (Matches

46;

Conservative

0

<u>..</u> Length Indels

, •

Gaps

0

Similarity

4.3%;

Score 43; Pred. No. Mismatches

DB 1;

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The present invention relates to oligonucleotides encoding polymorphic convariants of proteins related to amylases, amyloid proteins, angiopoietin, apoptosis related proteins, cadherin, cyclin, polymerase, oncogenes, chistones, kinases, colony stimulating factors, complement related proteins, cytochromes, kinesins, cytokines, interferons, interfevens, interfevens, conceptions and thosesterases. The present sequence is one conception of the prevention, diagnosis and treatment of diseases consecuted with inappropriate expression of the proteins listed above. Disorders that may be prevented, diagnosis and treatment of diseases consecuted with inappropriate expression of the proteins listed above. Disorders that may be prevented, diagnosed and/or treated include consecutes (e.g. rheumatoid arthritis, multiple sclerosis, diabetes, consecuted consecuted by the proteins and crave's disease), inflammation, cancer (e.g. cancers of the bladder, brain, breast, colon and kidney, concerning the component consecutions of the proteins of pathogenic consecutions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Polymorphic nucleic acids encoding e.g. amylases, cyclins, polymerases, oncogenes and histones, useful for diagnosing and treating, e.g. cancer autoimmune diseases and infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Shimkets RA,
Sequence 51 BP; 10 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; Page 2678; 4143pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2001-465210/50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Leach M;
    15
    Ç,
    17 G;
    9 T;
       0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    e.g. cancer,
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RESULT 70 AAL31458 Immunosuppressive; immunostimulatory; antiinflammatory; cytostatic; neuroprotective; antimicrobial; gene therapy; vaccine; amylase; cancer amyloid protein; angiopoietin; apoptosis related protein; cadherin; cyclin; polymerase; oncogene; histone; kinase; colony stimulating fact; complement related protein; cytochrome; kinesin; cytokine; interferon; interleukin; G-protein coupled receptor; thioesterase; inflammation; multifactorial disease; autoimmune disease; infection; nervous system disease; ss. Homo sapiens Human SNP oligonucleotide #4666. 24-JAN-2002 AAL31458 standard; DNA; 51 845 TGCCTCGGCCTCCCAAAGTGCTGGGATTACAGGCGTGAGCCACCACGCCCG 895 51 TGCCTTGGCCTCCCAAAGTGTTGGGGTTACAGGCGTGAGCCACCACCCCAG 1 (first entry) ВP cancer;

DE XXX RR XX R

WPI; 2001-465210/50

Shimkets RA,

Leach M;

(CURA-) CURAGEN CORP

28-DEC-1999; 27-DEC-2000;

99US-0173419P 2000US-00173419

28-DEC-2000; 2000WO-US035498

05-JUL-2001. WO200147944-A2

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RESULT 71
AAL31460
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention relates to oligonucleotides encoding polymorphic variants of proteins related to amylases, amyloid proteins, angiopoietin, angiopoietin, appotosis related proteins, cadherin, cyclin, polymerase, oncogenes, chistones, kinases, colony stimulating factors, complement related proteins, cytochromes, kinesins, cytokines, interferons, interleukins. Grotein coupled receptors and thioesterases. The present sequence is one such oligonucleotide. The oligonucleotides and the peptides encoded by them may be used in the prevention, diagnosis and treatment of diseases cassociated with inappropriate expression of the proteins listed above. Disorders that may be prevented, diagnosed and/or treated include consorters that may be prevented, diagnosed and/or treated include diseases (e.g. rheumatoid arthritis, multiple sclerosis, diabetes, systemic lupus erythromatosus and Grave's disease), inflammation, cancer (e.g. cancers of the bladder, brain, breast, colon and kidney, leukaemia), diseases of the nervous system and an infection of pathogenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
Polymorphic nucleic acids encoding e.g. amylases, cyclins, polymerases, oncogenes and histones, useful for diagnosing and treating, e.g. cancer
                                                                                                                                                                28-DEC-1999;
27-DEC-2000;
                                                                                                                                                                                                                                                                                                                                                                              Immunosuppressive; immunostimulatory; antiinflammatory; cytostatic; neuroprotective; antimicrobial; gene therapy; vaccine; amylase; cancer amyloid protein; angiopoietin; apoptosis related protein; cadherin; cyclin; polymerase; oncogene; histone; kinase; colony stimulating fact; complement related protein; cytochrome; kinesin; cytokine; interferon; interleukin; G-protein coupled receptor; thioesterase; inflammation; multifactorial disease; autoimmune disease; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human SNP oligonucleotide #4668.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAL31460 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Polymorphic nucleic acids encoding e.g. amylases, cyclins, polymerases, oncogenes and histones, useful for diagnosing and treating, e.g. cancer, autoimmune diseases and infections.
                                                        WPI; 2001-465210/50
                                                                                                                                                                                                                    28-DEC-2000; 2000WO-US035498
                                                                                                                                                                                                                                                                                          WO200147944-A2
                                                                                                                                                                                                                                                                                                                           Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                               nervous system
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1008
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                                                                                                                                                                99US-0173419P
2000US-00173419
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                         Leach
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      51
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 43;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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 e.g. cancer,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cancer;
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RESULT 72
AAI77523/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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Best Local
AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide sequences (I), which contain single nucleotide polymorphisms (SNPs). AAM53114 to AAM53329 represent peptides related to human polymorphic polynucleotide sequences. The sequences can be used in gene and protein therapy, and in vaccine production. (I) and the polypeptides encoded by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; single nucleotide polymorphism; SNP; genome; gene ther protein therapy; vaccine; probe; diagnostic assay; detection; quantitation; restorative therapy; polymorphic; ds.
                                                                                                                                                                                                                                                                                                                30-NOV-1999;
29-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                       30-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human silent SNP containing nucleic acid SEQ:4464
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 51 BP; 10 A; 16 C; 14 G; 11 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 autoimmune diseases and infections
                                                                                                                                                                   Polymorphic nucleic acid sequences, useful in genetic testing
                                                                                                                                                                                                       WPI; 2001-356160/37
                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200140521-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAI77523
                                                                                                                                                                                                                                                                               (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       849 TCGGCCTCCCAAAGTGCTGGGATTACAGGCGTGAGCCACCACCACCGCCTT 899
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        46;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        standard;
                                                                                                              Page 1877; 2653pp; English
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Pred. No. 1.
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RESULT 73
AAI78387
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Best Local S
Matches 46
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AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide sequences (I), which contain single nucleotide polymorphisms (SNPs). AAM53114 to AAM53129 represent peptides related to human polymorphic polynucleotide sequences. The sequences can be used in gene and protein therapy, and in vaccine production. (I) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of polymorphic polypeptides. For example, (I) may be used to treat disorders by rectifying mutations or deletions in a patient's genome that affect the activity of polypeptides by expressing inactive proteins or to supplement the patients own production of polypeptide. Additionally, (I) and its complementary
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                                                                                                                                                                                                                                                                                                                                                                  Claim 1; Page 2140; 2653pp; English
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RESULT 74
AAI76192/c
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AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide Sequences (1), which contain single nucleotide polymorphisms (SNPs). CC AAM53114 to AAM53329 represent peptides related to human polymorphic cypolynucleotide sequences. The sequences can be used in gene and protein cypolynucleotide sequences. The sequences can be used in gene and protein cypolynucleotide sequences. The sequences can be used in gene and protein cypolynucleotide with inappropriate expression of polymorphic polypeptides. For company, and in vaccine production, diagnosis and treatment of diseases company, and in a patient's genome that affect the activity of polypeptides. For company, and in a patient's genome that affect the activity of polypeptides by expressing inactive proteins or to supplement the patients own componentary company also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and company also be in need of restorative therapy. The conjugation of polypeptides be used as antigens in the production of conjugation and activity. The antibodies may also be used to down regulate expression and activity. The antibodies may also
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29-NOV-2000; 2000US-00726173
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AAM53114 to AAM53329 represent peptides related to human polymorphic polymucleotide sequences. The sequences can be used in gene and protein therapy, and in vaccine production. (I) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of polymorphic polypeptides. For example, (I) may be used to treat disorders by rectifying mutations or deletions in a patient's genome that affect the activity of polypeptides
                            by expressing inactive proteins or to supplement the patients own production of polypeptide. Additionally, (I) and its complementary sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and therefore which patients may be in need of restorative therapy. The polypeptides encoded by (I) may be used as antigens in the production of antibodies specific for polymorphic polypeptides. The antibodies may also be used to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic polypeptides in samples
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29-NOV-2000; 2000US-00726173.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; single nucleotide polymorphism; SNP; genome; protein therapy; vaccine; probe; diagnostic assay; quantitation; restorative therapy; polymorphic; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human silent SNP containing nucleic acid SEQ:4524.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAI77583;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAI77583 standard; DNA; 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         be used as diagnostic agents for detecting the presence of polymorphic polypeptides in samples
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local
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   51
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 BP;
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ilarity 90.2%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP; 14 A;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Leach M,
 15
 A; 7 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     뫄
 20 G; 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 43; DB:
Pred. No. 1.7e.
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             9 T;
 7,
 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0 U; 0 Other;
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ď,
 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      gene therapy;
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Query Match

Score 43;

DB

1; Length 51;

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RESULT 76
RAIT3060/c
ID AAIT30
XX
AAIT3060/c
ID AAIT30
XX
AAIT30
XX
XX
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Human
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AAIT31
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PI Casmoli
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PI Casmoli
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AAIT31

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                                                                                                                                                                                                                                                                                                             CC sequences (1), which contain single nucleotide polymorphisms (SNPs).

CC AAM53114 to AAM53329 represent peptides related to human polymorphic CC polymucleotide sequences. The sequences can be used in gene and protein CC thermapy, and in vaccine production. (1) and the polymorphism cassociated with inappropriate expression of polymorphic polymorphic cc example, (1) may be used to treat disorders by rectifying mutations or CC example, (1) may be used to treat disorders by rectifying mutations or CC production of polypeptides. Additionally, (1) and its complementary CC sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and therefore which patients may be in need of restorative therapy. The CC polypeptides encoded by (1) may be used as antigens in the production of contibodies specific for polymorphic polypeptides. The antibodies may also be used as diagnostic agents for detecting the presence of polypeptides in samples and cc polypeptides in samples as antibodies may also be used as diagnostic agents for detecting the presence of polymorphic polypeptides. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic polypeptides in samples in samples in samples in samples in samples in samples in samples.
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                                                                                                                              Query Match
Best Local Similarity
Matches 46; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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                                                                                                                                                                                                                                                             Sequence 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide sequences (I), which contain single nucleotide polymorphisms (SNPs).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1; Page 54; 2653pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Polymorphic nucleic acid sequences, useful in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-356160/37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Shimkets RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30-NOV-1999; 99US-0168138P.
29-NOV-2000; 2000US-00726173.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (CURA-) CURAGEN CORP.
                                         684 CCTCTGCCCCGGGTTCAAGTTATTCTCCTGCCCAGCCTCCTGAGTAGC 734
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    681
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   CCTCCGCCTCCTGGGTTCAAGCGATTCTCCTGCCTCAGCCTCCTGAGTAGC 1
                                                                                                                                                                                                                                                             BP; 13 A; 11 C; 21 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
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                                                                                                                                  Conservative
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0; Mismatches 5;
                                                                                                                                                             Score 43; I
Pred. No. 1.
                                                                                                                                  Mismatches
                                                                                  DB 1;
1.7e+02;
5; Indels
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RESULT 77
AAI79782/c
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AAI73863
ID AAI73
XX
                                                                                                                                                                                                                                                                              AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide sequences (I), which contain single nucleotide polymorphisms (SNPs). CC AAM53114 to AAM5329 represent peptides related to human polymorphic polynucleotide sequences. The sequences can be used in gene and protein therapy, and in vaccine production. (I) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases company, and the interpretations of polymorphic polypeptides. For company, and the prevention, diagnosis and treatment of diseases company, and the prevention, diagnosis and treatment of diseases company, and the production of diseases company, and the prevention of polypeptides. For completions in a patient's genome that affect the activity of polypeptides. For company and the presence of genoment the patients own company inactive proteins or to supplement the patients own company. (I) and its complementary company also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and therefore which patients may be in need of restorative therapy. The company production of antibodies specific for polymorphic polypeptides. The antibodies may also be used to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic polypeptides in samples.
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                                                                                                                                                                             Best Loc
Matches
                                                                                                                                                                                                                Query Match
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AAI79782 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       30-NOV-1999; 99US-0168138P
29-NOV-2000; 2000US-00726173
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 1; Page 2562; 2653pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Polymorphic nucleic acid sequences, useful in genetic testing
                                                                                                                                                                                                                                                  Sequence 51
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                                                                                                                                                                             Local Similarity 90.2
nes 46; Conservative
                                                                                                                                           989
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                                                                                                                               GCCTCCCGGGCTCAAGCGATTCTCCTGTCTCAGCCTCCCAAGCAGCAGCTGGGA 1039
                                                                                                             GCCTCCTGGGTTCAAGCAATTCTCCTGCCTCAGCCTCCCAAGTAGCTGGGA 1
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                                                                                                                                                                                                                                                  A; 12 C; 18 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                             4.3%;
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                                                                                                                                                                             Score 43; DB
Pred. No. 1.7e
0; Mismatches
                                                                                                                                                                              0,
                                                                                                                                                                         Db _
1.7e+02;
5;
                                                                                                                                                                                                                Length 51;
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                                                                                                                                                                              0; Gaps
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AAI73863 standard; DNA; 51

ВÞ

Human silent SNP containing nucleic acid SEQ:4523

09-NOV-2001

(first entry)

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CC AAM53114 to AAM53329 represent peptides related to human polymorphic polymorphic conduction to polymorphic conduction therapy, and in vaccine production. (I) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases conducted with inappropriate expression of polymorphic polypeptides. For considered with inappropriate expression of polymorphic polypeptides. For conductions in a patient's genome that affect the activity of polypeptides or conduction of polypeptide. Additionally, (I) and its complementary conduction of polypeptide. Additionally, (I) and its complementary conduction of polypeptide. Additionally, (I) and its complementary conduction patients may be in need of restorative therapy. The colypeptides encoded by (I) may be used as antigens in the production of polypeptides for nolumerative therapy. The polypeptides encoded by (I) may be used as antigens in the production of contribution of contributions of contributions of contribution of contributions of contributi
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       antibodies specific for polymorphic polypeptides. The antibodies may also be used to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic
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29-NOV-2000; 2000US-00726173
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 51 BP;
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0; Mismatches
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Pred. No. 1.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            T; 0 U; 0 Other;
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RESULT 80
AAI73063/C
ID AAI730
XX AAI730
AC AAI730
XX 09-NOV
XX Uprotei
KW protei
KW quanti
XX
OS Homo s
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Best Local Similarity
Matches 46; Conserv
                      Homo sapiens
                                                          Human; single nucleotide polymorphism; SNP; genome; gene therapy; protein therapy; vaccine; probe; diagnostic assay; detection; quantitation; restorative therapy; polymorphic; ds.
                                                                                                                                        Human silent
                                                                                                                                                                                09-NOV-2001
                                                                                                                                                                                                                                                             AAI73063 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 51 BP; 15
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29-NOV-2000; 2000US-00726173
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 polypeptides in samples
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                                                                                                                                                                                                                                                                                                                                                                                                    681
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                                                                                                                                           SNP
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                                                                                                                                      containing nucleic acid SEQ:4.
                                                                                                                                                                                                                                                             DNA;
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                                                                                                                                                                                entry)
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                                                                                                                                                                                                                                                                 ВP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 43;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                           <u>5</u>
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RESULT 81
AAI76093/c
ID AAI760
XX AAI760
XX AAI760
XX AAI760
XX AAI760
XX Human
XX Human;
KW Protei
KW Quanti
XX Quanti
XX WO2001
XX WO2001
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XX WO2001
XX Y
PD 07-JUN
XX Y
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAI73060 to AAI79867 represent isolated human polymorphic polymucleotide sequences (I), which contain single nucleotide polymorphisms (SNPs).

CC AAM53114 to AAM53329 represent peptides related to human polymorphic polymorphic CC polymucleotide sequences. The sequences can be used in gene and protein CC therapy, and in vaccine production. (I) and the polypeptides encoded by CC them may be used in the prevention, diagnosis and treatment of diseases CC associated with inappropriate expression of polymorphic polypeptides. For CC example, (I) may be used to treat disorders by rectifying mutations or CC deletions in a patient's genome that affect the activity of polypeptides by expressing inactive proteins or to supplement the patients own CC production of polypeptide. Additionally, (I) and its complementary CC sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and CC polypeptides encoded by (I) may be used as antigens in the production of CC polypeptides encoded by (I) may be used as antigens in the production of CC polypeptides specific for polymorphic polypeptides. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic colypeptides may also be used as diagnostic agents for detecting the presence of polymorphic polypeptides may also be used as diagnostic agents for detecting the presence of polymorphic colypeptides may also be used as diagnostic agents for detecting the presence of polymorphic polypeptides in samples may also be used as diagnostic agents for detecting the presence of polymorphic polypeptides may also be used as diagnostic agents for detecting the presence of polymorphic polypeptides may also be used as diagnostic agents for detecting the presence of polymorphic polypeptides.
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                                                                                                                                                                                                    Human; single nucleotide polymorphism; SNP; genome; protein therapy; vaccine; probe; diagnostic assay; oquantitation; restorative therapy; polymorphic; ds.
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                                                                                                                                                                                                                                                                                                                                                                                   AAI76093;
                                                                                                                                                                                                                                                                                                                                                                                                                               AAI76093 standard; DNA; 51.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 51 BP; 12 A; 10 C;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Polymorphic nucleic acid sequences, useful in genetic testing and
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29-NOV-2000; 2000US-00726173
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                                                                   07-JUN-2001.
                                                                                                             WO200140521-A2
                                                                                                                                                                                                                                                                                        Human silent SNP containing nucleic acid SEQ:3034.
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                                                                                                                                                                                                                                                                                                                                                                                                                                  ВP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 43; DB 1; Lo
Pred. No. 1.7e+02;
0; Mismatches 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 51
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                                                                                                                                                                                                                             detection
                                                                                                                                                                                                                                                gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      <u>,</u>
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30-NOV-2000; 2000WO-US032758

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RESULT 82
AAI76247/c
D X A X A X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X 
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Matches 46
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29-NOV-2000; 2000US-00726173.
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Shimkets
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29-NOV-2000; 2000US-00726173.
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                                                            (CURA-) CURAGEN CORP.
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Pred. No. 1.7e
0; Mismatches
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polypeptides in samples
                                                      be used as diagnostic agents for detecting the presence of polymorphic
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문 Query Match Best Local S Matches 46 697 GGTTCAAGTTATTCTCCTGCCCCAGCCTCCTGAGTAGCTGGGACTACAGGC 747 51 46; Similarity derrenadecantretectecercadectecdadradeteddadeadeadec 1 Conservative 90.2%; 0 Score 43; I Pred. No. 1. Mismatches JB 1, 1.7e+02; 5; DB 1; Indels <u>,,</u> Gaps

0

Sequence 51 BP; 10 A; 14 C; 18 G; 9 T; 0 U; 0 Other;

RESULT 83
AAI78389
ID AAI78 Human; single nucleotide polymorphism; SNP; genome; protein therapy; vaccine; probe; diagnostic assay; c quantitation; restorative therapy; polymorphic; ds. Human silent SNP containing nucleic acid SEQ:5330. AAI78389; AAI78389 standard; 09-NOV-2001 (first entry) DNA; 51 ВP detection;

gene therapy;

Homo sapiens.

WO200140521-A2

30-NOV-2000; 2000WO-US032758 07-JUN-2001

30-NOV-1999; 99US-0168138P 29-NOV-2000; 2000US-00726173

(CURA-) CURAGEN CORP.

Shimkets RA, Leach M;

WPI; 2001-356160/37

Polymorphic nucleic acid sequences, useful in genetic testing and therapy.

Claim 1; Page 2141;

2653pp;

English.

to

AAI79867

Query Match

Local Similarity

4.3%;

Score 43; Pred. No.

1.7e+02; DB 1;

Length 51;

Sequence 51 BP; 10 A; 20 C;

13 G; 8 T; 0 U; 0 Other;

CC therapy, and in vaccine production. (1) and the polypeptides encoded by CC them may be used in the prevention, diagnosis and treatment of diseases CC associated with inappropriate expression of polymorphic polypeptides. For CC example, (I) may be used to treat disorders by rectifying mutations or CC example, (I) may be used to treat disorders by rectifying mutations or CC example, (I) may be used to treat disorders by rectifying mutations or CC by expressing inactive proteins or to supplement the patients own CC production of polypeptide. Additionally, (I) and its complementary CC sequences may also be used as DNA probes in diagnostic assays to detect CC and quantitate the presence of similar nucleic acids in samples, and CC therefore which patients may be in need of restorative therapy. The CC polypeptides encoded by (I) may be used as antigens in the production of CC antibodies specific for polymorphic polypeptides. The antibodies may also be used to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic CC polypeptides in samples

polynucleotide sequences. The sequences can be used in gene and protein AAM53114 to AAM53329 represent peptides related to human polymorphic

represent isolated human polymorphic polynucleotide contain single nucleotide polymorphisms (SNPs).

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RESULT 84
AAH89507
ID AAH89
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AAH89507
AC AAH82
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The present invention relates to human nucleic acids containing single nucleotide polymorphisms (SNPs). These can be used in forensic and paternity tests, and to aid in the treatment of diseases associated with aberrant protein expression, including cancer, amyloidosis, diabetes,
                                                                                                                                                                                                                                                                                                       Isolated human polynucleotides containing single nucleotide polymorphisms, useful for the treatment and diagnosis of e.g.
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Matches Query Match

46;

Conservative

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4.3%; 90.2%; 13 C;

Score 43; D Pred. No. 1. Mismatches

DB 1; .7e+02;

Length 51;

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Local Similarity

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Best Local
                             The present invention relates to human nucleic acids containing single nucleotide polymorphisms (SNPs). These can be used in forensic and paternity tests, and to aid in the treatment of diseases associated with aberrant protein expression, including cancer, amyloidosis, diabetes, Alzheimer's disease, Down's syndrome, oedema, lupus (SLB), vasculitis, glomerulonephritis, haemolytic anaemia, thrombocytopaenia, arthritis, meningitis, muscular disporders, dementia, neurological diseases, tuberous sclerosis, male infertility, hypercalcaemia, blood pressure disorders, osteoporosis, pathogenic infections, hypercholesterolaemia, obesity or autoimmunity. The present sequence is a polymorphism-containing oligonucleotide fragment of the invention
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RESULT 86
AAH89404
RESULT 87
AAH89466/c
ID AAH894
XX
AC AAH894
XX
DT 01-OCT
XX
DE Human
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                                                                                                                                                       밁
                                                                                                                                                                                  Ś
                                                                                                                                                                                                             Query Match
Best Local S
Matches 46
                                                                                                                                                                                                                                                                                            The present invention relates to human nucleic acids containing single nucleotide polymorphisms (SNPs). These can be used in forensic and paternity tests, and to aid in the treatment of diseases associated with aberrant protein expression, including cancer, amyloidosis, diabetes, Alzheimer's disease, Down's syndrome, oedema, lupus (SLE), vasculitis, glomerulonephritis, haemolytic anasemia, thrombocytopaenia, arthritis, meningitis, muscular disorders, dementia, neurological diseases, tuberous sclerosis, male inferrility, hyperclacemia, blood pressure disorders, osteoporosis, pathogenic infections, hypercholesterolaemia, obesity or autoimmunity. The present sequence is a polymorphism-containing oligonucleotide fragment of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human coding sequence polymorphic site SEQ ID NO: 185
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAH89404 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        07-JAN-2000; 2000US-0174962P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    05-JAN-2001; 2001WO-US000322
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19-JUL-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  polymorphisms, infection and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2001-451871/48.
P-PSDB; AAM00291.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200151670-A2
                                                                                                                                                                                                                                                                      Sequence 51 BP; 9 A; 20 C; 9 G; 13 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1; Page 159; 475pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  rsolated human polynucleotides containing single nucleotide
polymorphisms, useful for the treatment and diagnosis of e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     forensic test;
                              01-OCT-2001
                                                                                     AAH89466 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (CURA-) CURAGEN CORP.
                                                                                                                                                                                    663
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  single nucleotide polymorphism; SNP;
                                                                                                                                                         μ
                                                                                                                                                                                                              46;
                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AGTAGAGACGGGGTTTCACCATGTTAGCCAGGCTGGTCTCAAACTCCTGAC 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTAGAGGCGGGGTTTTCACCATATTTGTCAGGCTGGTCTCAAACTCCTGAC 1133
                                                                                                                                                                         CGCAATCTTGGCTCACTGCAACCTCTGCCTCCCGGGTTCAAGTTATTCTCC 713
                                                                                                                                                         CACGATCTTGGCTCACTGCAACCTCTGCCTCCCAGGTTCAAGTGATCCTCC 51
                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
 sequence polymorphic site SEQ ID NO: 247.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Leach
                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   diabetes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     aberrant protein expression;
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                                                                                     DNA;
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                                                                                                                                                                                                                           90.2%;
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                                                                                     ВP
                                                                                                                                                                                                             Score 43; DB 1; ~~
Pred. No. 1.7e+02;
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                                                                                                                                                                                                                                          Length 51;
                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cancer,
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RESULT 88
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Best Local &
                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to human nucleic acids containing single nucleotide polymorphisms (SNPs). These can be used in forensic and paternity tests, and to aid in the treatment of diseases associated with aberrant protein expression, including cancer, amyloidosis, diabetes, Alzheimer's disease, Down's syndrome, oedema, lupus (SLE), vasculitis, glomerulonephritis, haemolytic anaemia, thrombocytopaemia, arthritis, meningitis, muscular discorders, dementia, neurological diseases, tuberous sclerosis, male inferrility, hypercalcaemia, blood pressure disorders, osteoporosis, pathogenic infections, hypercholesterolaemia, obesity or autoimmunity. The present sequence is a polymorphism-containing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; single nucleotide polymorphism; SNP; forensic test; aberrant protein expression;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       05-JAN-2001; 2001WO-US000322
                                                                                                                                                                                                                                                                                                                                                                            Sequence 51 BP; 13 A; 11 C; 21 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1; Page 175; 475pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Shimkets RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      07-JAN-2000; 2000US-0174962P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200151670-A2
                                                                                                                                                                                                                                                                                                                                                                                                                              oligonucleotide fragment of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Isolated human polynucleotides containing single nucleotide polymorphisms, useful for the treatment and diagnosis of e.c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         P-PSDB; AAM00351.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19-JUL-2001.
AAH39240;
                                            AAH39240 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     infection and diabetes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2001-451871/48.
                                                                                                                                                                                                    817 TCTTGATCTCTGGACCTTGTGATCTGCCTGCCTCGGCCTCCCAAAGTGCTG 867
                                                                                                                                                                            51
                                                                                                                                                                                                                                                                           46;
                                                                                                                                                                                                                                                                                                     Similarity
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                                                                                                                                                                                                                                                                                Conservative
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                                                                                                                                                                                                                                                                                                  4.3%;
                                                 51
                                                                                                                                                                                                                                                                                                                                                                                                                                     the invention
                                                                                                                                                                                                                                                                                                        Pred.
                                                                                                                                                                                                                                                                                                                             Score 43;
                                                                                                                                                                                                                                                                                Mismatches
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                                                                                                                                                                                                                                                                                                     1.7e+02;
                                                                                                                                                                                                                                                                                                                                  DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 paternity test; ds.
                                                                                                                                                                                                                                                                                                                               Length 51;
                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  of e.g. cancer,
                                                                                                                                                                                                                                                                                0,
                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         tuberous
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Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis; inflammation; forensic investigation; paternity analysis; ds.
                                                                                                                                                                                                                         Human SNP flanking oligonucleotide SEQ ID 2036.
                                                                                                                                                                                                                                                          14-AUG-2001
                                                                              Homo
                                                                              sapiens
                                                                                                                                                                                                                                                          (first entry)
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Human coding

26-APR-2001

WO200129262-A2

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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             osteogenesis imperfecta and acute intermittent porphyria. Phenotypic traits also include symptoms of or susceptibility to multifactorial disease of which a component is or may be genetic such as autoimmune disease, including, rheumatoid arthritis, multiple sclerosis, inflammation, cancer, nervous system diseases and infection by pathogenic microorganism. The method is also useful in forensic investigations and paternity analysis. The present sequence represents a fragment of human DNA flanking the site of a single nucleotide polymorphism
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SNP flanking sequence, the SNPE primer is used as a genotyping primer. The oligonucleotides are useful for genotyping anucleic acid sample by performing a single-nucleotide primer extension reaction. The performing a single-nucleotide primer extension reaction. The oligonucleotides are useful for determining the presence, absence or identity of a SNP and for genotyping nucleic acid samples, for e.g. to assess by association analysis the genotype of an individual or group c ansees by association analysis the genotype of an individual or group c individuals, having a pathological phenotypic trait suspected of being caused by one or more SNPs. Phenotypic traits include diseases e.g. agammaglobulinaemia, diabetes insipidus, Lesch-Nyhan syndrome, muscular dystrophy, familial hypercholesterolaemia, polycystic kidney disease,
22-NOV-2000; 2000WO-US032311.
                                                                            WO200138586-A2
                                                                                                                                                                          immunosuppressive; antiinflammatory; neuroprotective; antimicrobiautoimmune disease; inflammation; cancer; nervous system disease;
                                                                                                                                                                                                                   Human; single nucleotide polymorphism; SNP; polymorphism;
                                                                                                                                                                                                                                                           Human silent noncoding SNP oligonucleotide SEQ ID NO:114.
                                                                                                                                                                                                                                                                                                  05-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                 ABL00123 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 51 BP; 7 A; 17 C; 16 G; 11 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              includes kits for determining the presence or absence of a SNP, using the oligonucleotides of the invention. The PCR primers are used to amplify a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     primer extension (SNPE) primers, and the sequences of regions flanking sites of single nucleotide polymorphisms SNPs. The present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequences AAH37205 -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; Page 60; 83pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         acid sample.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-290930/30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Picoult-Newburg L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13-OCT-2000; 2000WO-US028436
                                                                                                                                                       infection; polymorphic protein; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         646 AGGCTGGAGTGCAGTGGCGCAATCTTGGCTCACTGCAACCTCTGCCTCCCG 696
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               46;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORCHID BIOSCIENCES INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AGGCTGGGGTGCAGTGCGATCTCGGCTCACTGCAACCTCTACCTCCCG 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            99US-0160096P.
                                                                                                                                                                                                                                                                                                                                                                                 DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAH40944 represent PCR primers, single nucleotide
                                                                                                                                                                                                                                                                                                                                                                                 51
                                                                                                                                                                                                                                                                                                                                                                                 ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 43; DB Pred. No. 1.7e 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    .7e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ټ.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                   cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ٥,
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                oligonucleotides. The sequences from the present invention can have immunosuppressive, cytostatic, antiinflammatory, neuroprotective and antimicrobial activities. Nucleic acids, polypeptides, oligonucleotides and antibodies from the present invention can be used for treating a subject suffering from, at risk for, or suspected of, suffering from a pathology ascribed to the presence of a sequence polymorphism. The pathology may be autoimmune diseases, inflammation, cancer, diseases of the nervous system, and infection by pathogenic microorganisms. The SNPs are also useful for determining which forms of a characterised polymorphism are present in individuals. The antibodies may be used in the detection, quantitation and/or cellular or tissue localisation of a
                                                                                                                                                                                                                                                                                                          Human; single nucleotide polymorphism; SNP; polymorphism; cytostatic; immunosuppressive; antiinflammatory; neuroprotective; antimicrobial; autoimmune disease; inflammation; cancer; nervous system disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          the detection, quantitation and/or cellular or tissue localisation of a polymorphic protein (e.g., for use in measuring levels of the polymorphic protein within appropriate physiological samples)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABL00010 to ABL01104 represent human nucleic acid oligonucleotides comprising one or more single nucleotide polymorphisms (SNPs). ABB to ABB56903 represent human peptides encoded by some of the SNP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Isolated human nucleic acids comprising one or more single nucleotide polymorphisms, useful for treating a subject suffering from a pathology, e.g. autoimmune diseases, ascribed to the presence of a sequence
                                                                                                                                                                                                              WO200138586-A2
                                                                                                                                                                                                                                                                                                                                                                                      Human silent noncoding SNP oligonucleotide SEQ ID NO:12.
                                                                                                                                                                                                                                                                                                                                                                                                                            05-MAR-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABL00021 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 51 BP; 16 A; 14 C; 11 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; Page 280; 674pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2001-355949/37.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              24-NOV-1999;
                                                                                                                                  22-NOV-2000; 2000WO-US032311.
                                                                                                                                                                        31-MAY-2001.
                                                                                                                                                                                                                                                                                      infection; polymorphic protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (CURA-) CURAGEN CORP.
                                                          (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          181 TAGAGATGGAGTTTCTCCATGTTGGTCAGGCTGGTCTCGAACTCCCGACCT 231
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                46;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                99US-0167383P
                                                                                              99US-0167383P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 43; DB 1;
Pred. No. 1.7e+02;
                                                                                                                                                                                                                                                                                            dg.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 51;
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Shimkets RA,

Leach M;

Claim 1;

Page 248; 674pp; English.

ABL00010

e.g. autoimmune polymorphism.

WPI; 2001-355949/37

Isolated human nucleic acids comprising one or more single nucleotide polymorphisms, useful for treating a subject suffering from a pathology, e.g. autoimmune diseases, ascribed to the presence of a sequence

comprising one or more single nucleotide polymorphisms (SNPs). ABB56 to ABB56903 represent human peptides encoded by some of the SNP oligonucleotides. The sequences from the present invention can have

to ABL01104 represent human nucleic acid oligonucleotides

ABB56531

imminosuppressive, cytostatic, antiinflammatory, neuroprotective and antimicrobial activities. Nucleic acids, polypeptides, oligonucleotides and antibodies from the present invention can be used for treating a subject suffering from, at risk for, or suspected of, suffering from a pathology ascribed to the presence of a sequence polymorphism. The pathology may be autoimmune diseases, inflammation, cancer, diseases of the nervous system, and infection by pathogenic microorganisms. The SNPs are also useful for determining which forms of a characterised polymorphism are present in individuals. The antibodies may be used in the detection, quantitation and/or cellular or tissue localisation of a polymorphic protein (e.g., for use in measuring levels of the polymorphic

Sequence

51

BP; 7

A.

13 G; 11 T;

0 U; 0 Other;

Conservative

4.3%; 20 C;

DB 1; 1.7e+02;

Length 51 Indels

<u>5</u>

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Gaps

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protein within appropriate physiological samples)

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Best Local S
Matches 46
                                                                                                                                                                                                                                  Human; single nucleotide polymorphism; SNP; genome; gene therapy; protein therapy; vaccine; probe; diagnostic assay; detection; quantitation; restorative therapy; polymorphic; ds.
                                                                                                                                                                                                                                                                          Human silent SNP containing nucleic acid SEQ:3758
                                                                                                                      30-NOV-1999; 99US-0168138P
29-NOV-2000; 2000US-00726173.
                                                                                                                                                                                                                                                                                                 09-NOV-2001
                                                                                                                                                                                                                                                                                                                                        AAI76817 standard;
          Claim 1; Page 1201; 2653pp; English
                                                             WPI; 2001-356160/37
                                                                                                                                                   30-NOV-2000; 2000WO-US032758
                                                                                                                                                                         07-JUN-2001
                                                                                                                                                                                            WO200140521-A2
                                         Polymorphic nucleic acid sequences,
                                                                                                                                                                                                                                                                                                                                                                                                            987
                                                                                                                                                                                                                                                                                                                                                                                                                                1 Similarity
46; Conserv
                                                                                                   CURAGEN CORP
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                                                                                                                                                                                                                                                                                                 (first
                                                                               Leach M;
                                                                                                                                                                                                                                                                                                                                        DNA; 51
                                                                                                                                                                                                                                                                                                 entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                Score 43; DB
Pred. No. 1.7e
0; Mismatches
                                          useful
                                           ä
                                         genetic testing
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cc sequences (i), which contain single nucleotide polymorphic polynucleotide (i), which contain single nucleotide polymorphisms (SNPs).

Cc AAM53114 to AAM5329 represent peptides related to human polymorphic polynucleotide sequences. The sequences can be used in gene and protein therapy, and in vaccine production. (I) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases cascotated with inappropriate expression of polymorphic polypeptides. For cexample, (I) may be used to treat disorders by rectifying mutations or cc deletions in a patient's genome that affect the activity of polypeptides by expressing inactive proteins or to supplement the patients own cc production of polypeptide. Additionally, (I) and its complementary cc sequences may also be used as Dah probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and ct therefore which patients may be used of restorative therapy. The polypeptides encoded by (I) may be used as antigens in the production of the used to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic polypeptides. polypeptides in samples

Query Match Best Local (Matches Sequence 51 193 TTCTCCATGTTGGTCAGGCTGGTCTCGAACTCCCGACCTCAGATGATCC 45; Similarity TTCGCCATGTTGGCCAGGCTGGTCTCGAACTCCTGACCTCAGGTGATCC BP; 8 Conservative A; 17 C; 13 G; 13 4.3%; 0, Score 42.6 Pred. No. Mismatches 42.6; No. 1 T; 0 U; 0 Other; .8e+02 DB 1; Length Indels 49 241 0, Gaps

0

RESULT 92 ADI12541/c ID ADI125 Mutant human ADI12541; ADI12541 standard; 22-APR-2004 **BRCA1** (first entry) genomic DNA resulting DNA; 44 뫄 from deletion 3 SeqID

ovarian Homo sapiens cancer; recombination; mutant.

ds; cancer; human; tumour suppressor; breast cancer susceptibility gene 1; BRCA1; repetitive Alu.

WO2003104474-A2

18-DEC-2003.

09-JUN-2003; 2003WO-US018098

07-JUN-2002; 2002US-0387132P 09-AUG-2002; 2002US-0402430P

(MYRI-) MYRIAD GENETICS INC

Scholl T, Hendrickson BC, Ward В, Pruss

WPI; 2004-062369/06

Predicting a predisposition to cancer in a patient comprising detecting deletion in the BRCA1 gene that results from the unequal crossover between a pair of repetitive sequences in the BRCA1 gene.

Disclosure; SEQ ID NO 24; 59pp; English.

This invention relates to a novel method for predicting a predisposi to cancer in a patient by detecting large deletions in the human tum suppressor gene identified as the breast cancer susceptibility gene (BRCA1). Specifically, it refers to deletions that result from the unequal crossover between a pair of repetitive Alu sequences in the predisposition e human tumour in the BRCAI

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RESULT 93
AAH38364/c
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Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide primer extension (SNPE) primers, and the sequences of regions flanking sites of single nucleotide polymorphisms SNPs. The present invention includes kits for determining the presence or absence of a SNP, using the oligonucleotides of the invention. The PCR primers are used to amplify a SNP flanking sequence, the SNPE primer is used as a genotyping primer. The oligonucleotides are useful for genotyping a nucleit acid sample by performing a single-nucleotide primer extension reaction. The oligonucleotides are useful for determining the presence, absence or identity of a SNP and for genotyping nucleic acid samples, for e.g. to assess by association analysis the genotype of an individual or group of
                                                                                                                                                                                                                                                                                                                                                                                      Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2001-290930/30.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         13-OCT-2000; 2000WO-US028436.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SNPE; genotyping;
Lesch-Nyhan syndre
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14-AUG-2001
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                                                                                                                                                                                                                                                                                                                                                                                   55; 83pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               99US-0160096P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        disease; osteogenesis imperfecta; autoimmune disease; porphyria; rheumatoid arthritis; multiple sclerosis; porphyria; rheumatoid arthritis; multiple sclerosis; pare investigation; paternity analysis; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 oligonucleotide SEQ ID 1160
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA;
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97.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         51
                                                                                                                                                                                                                                                                                                                                                                                      English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pred. No. 1.60
); Mismatches
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                                                                                                                                                                         individuals, having a pathological phenotypic trait suspected of being caused by one or more SNPs. Phenotypic traits include diseases e.g. agammaglobulinaemia, diabetes insipidus, Lesch-Nyhan syndrome, muscular dystrophy, familial hypercholesterolaemia, polycystic kidney disease, osteogenesis imperfecta and acute intermittent porphyria. Phenotypic traits also include symptoms of or susceptibility to multifactorial disease of which a component is or may be genetic such as autoimmune disease, including, rheumatoid arthritis, multiple sclerosis, inflammation, cancer, nervous system diseases and infection by pathogenic microorganism. The method is also useful in forensic investigations and paternity analysis. The present sequence represents a fragment of human than the second of the present sequence represents a fragment of human than the second of the present sequence represents a fragment of human than the second of the second o
Sequence 51
                                                                                                                                flanking the site of a single nucleotide polymorphism
BP; 8
ð
14 C; 17 G; 12
      T, 0
      u; o
Other;
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Matches
                                                 Query Match
Best Local
                    854
51
                                       1 Similarity
43; Conserv
            CTCCCAAAGTGCTGGGATTACAGGCGTGAGCCACCACGCCCGGC
CTCCCAAAGTGCTGGGATTACAGGCGTGAGCCACCATGCCCGGC
                                       4.3%;
llarity 97.7%;
Conservative
                                        0
                                                  Score 42.4;
Pred. No. 1
                                         Mismatches
                                                  .8e+02;
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                                                            <u>ب</u>
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                                                            Length
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                    897
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RESULT 94
AAZ69411/
Human map-related biallelic marker SEQ ID
                                                                                               AAZ69411 standard; DNA;
                                 (first entry)
                                                                                                 47
   NO:3767
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Human genome; biallelic marker; high density disequilibrium map; genomic map; haplotype; phenotype; polymorphic base; genotyping; haplotyping; hybridisation; dentification; characterisation; distinct pingle nucleotide polymorphism; SNP; ds. diagnosis;

Homo sapiens

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28-OCT-1999
                                                           WO9954500-A2
                                                                    variation
                                                                 Location/Qualifiers replace(24,T)
/*tag= a
                                                               /standard_name= "single nucleotide
                                                               polymorphism"
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21-APR-1998; 23-NOV-1998; 21-APR-1999; (GEST) á GENSET Blumenfeld 98US-0082614P 98US-0109732P 99WO-IB000822 Z, Chumakov

qam Novel biallelic markers used genome. to construct a high density disequilibrium

2000-013267/01.

ω •• Page 1034; 2745pp; English.

AAZ65654 to AAZ69578 represent human biallelic markers from the pre invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AAZ69579 to AAZ77440 represent amplification primers for the biallelic markers. The biallelic markers of the inv have a variety of are useful in y of uses: they can be used for high density mapping of the and in complex association studies and haplotyping studies determining the genetic markers from the present position 24 of their invention

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RESULT 95
AAI74450/c
ID AAI744
CC sequences (I), which contain single nucleotide polymorphic polymucleotide CR sequences (I), which contain single nucleotide polymorphisms (SNPs). CC AAM53114 to AAM53329 represent peptides related to human polymorphic CR cham53114 to AAM53329 represent peptides related to human polymorphic CR cherapy, and in vaccine production. (I) and the polypeptides encoded by CR therapy, and in vaccine production. (I) and the polypeptides encoded by CR chem may be used in the prevention, diagnosis and treatment of diseases CR associated with inappropriate expression of polymorphic polypeptides. For CR cample, (I) may be used to treat disorders by rectifying mutations or CR cample, (I) may be used to treat disorders by rectifying mutations or CR captures in a patient's genome that affect the activity of polypeptides CR by expressing inactive proteins or to supplement the patients own CR production of polypeptide. Additionally, (I) and its complementary CR sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and CR polypeptides encoded by (I) may be used as antigens in the production of conjugations and conjugations of antibodies may also be used to down regulate expression and activity. The antibodies may also can be used to down regulate expression and activity. The antibodies may also
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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N.B. The SEQ ID NOS 2852, 2913, 2974, 3035, 3096, 3157, 3227, 3297 and 3367, are not actually given a sequence in the Sequence Listing from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; single nucleotide polymorphism; SNP; genome; protein therapy; vaccine; probe; diagnostic assay; quantitation; restorative therapy; polymorphic; ds.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    30-NOV-1999; 99US-0168138P
29-NOV-2000; 2000US-00726173
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 1; Page 479; 2653pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Polymorphic nucleic acid sequences, useful in genetic testing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-356160/37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Shimkets RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-NOV-2000; 2000WO-US032758
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       673 GCTCACTGCAACCTCTGCCTCCCGGGTTCAAGTTATTCTCCCTGCCCC 719
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Leach M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            containing nucleic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4.3%;
93.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          51
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Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 47,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
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XAXFXGX&&&&&&&&&&&&&X

Query Match

Sequence

51

BP;

15

A; 10 C;

18 G; 8 T; 0 U; 0 Other,

Score 42.2;

DB 1;

Length 51;

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RESULT 96
AAI76248/c
ID AAI762
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CR AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide consequences (I), which contain single nucleotide polymorphisms (SNPB).

CR AAM53114 to AAM5329 represent peptides related to human polymorphic consequences. The sequences can be used in gene and protein therapy, and in vaccine production. (I) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases cassociated with inappropriate expression of polypeptides. For example, (I) may be used to treat disorders by rectifying mutations or condections in a patient's genome that affect the activity of polypeptides by expressing inactive proteins or to supplement the patients own conduction of polypeptide. Additionally, (I) and its complementary consequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and therefore which patients may be in need of restorative therapy. The polypeptides encoded by (I) may be used as antigens in the production of antibodies specific for polymorphic polypeptides. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic polypeptides. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic polypeptides may also be used as diagnostic agents for detecting the presence of polymorphic polypeptides in samples.
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; single nucleotide polymorphism; SNP; genome, protein therapy; vaccine; probe; diagnostic assay; quantitation; restorative therapy; polymorphic; ds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 51 BP; 10 A; 11 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        09-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAI76248;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAI76248
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          be used as diagnostic agents for detecting the presence of polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30-NOV-2000; 2000WO-US032758
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    07-JUN-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human silent SNP containing nucleic acid SEQ:3189
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        polypeptides in
                                                                                                                                                                                                                                                                                                                                                               Claim 1; Page 1026; 2653pp;
                                                                                                                                                                                                                                                                                                                                                                                                                 Polymorphic nucleic acid sequences, useful in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   29-NOV-2000; 2000US-00726173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200140521-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (CURA-) CURAGEN CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    99US-0168138P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        samples
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         51
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                                                                                                                                                                                                                                                                                                                                                                 English.
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Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         G; 11 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          42.2;
No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          .9e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    genome;
                                                                                                                                                                                                                                                                                                                                                                                                                   genetic testing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            <u>,</u>
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AAI75601/c
ID AAI75601 standard; DNA; 51
                                                                                                                                                          CC sequences (I), which contain single nucleotide polymorphisms (SNPs).

CR AAM53114 to AAM5329 represent peptides related to human polymorphic may also be used to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic polymorphic.
                                                                     Query Match
Best Local :
                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2001-356160/37.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Shimkets
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30-NOV-1999; 99US-0168138P.
29-NOV-2000; 2000US-00726173.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-NOV-2000; 2000WO-US032758.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            07-JUN-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; single nucleotide polymorphism; SNP; genome; gene therapy; protein therapy; vaccine; probe; diagnostic assay; detection; quantitation; restorative therapy; polymorphic; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human silent SNP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            09-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                            AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide
                                                                                                                                                  polypeptides in samples
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (CURA-) CURAGEN CORP.
                                                                     Local Similarity
                            634
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48
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ₽,
               ACTCTGTCACCCAGGCTGGAGTGCAGTGGCGCAATCTTGGCTCACTG 680
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATCTTGGCTCACTGCAACCTCTGCCTCCTGGGTTCAAGCGATTCTCC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATCTTGGCTCACTGCAACCTCTGCCTCCCGGGTTCAAGTTATTCTCCC
ACTCTGTCGCCCAGGCTGGAGTACAGTGGCACAATCTTGGCTCACTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Page 829;
                                                       4.3%;
nilarity 93.6%;
Conservative
                                                                                                                     BP; 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       nucleic acid sequences, useful in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Leach
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             containing nucleic acid SEQ:2542
                                                                                                                     A; 13 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           2653pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         93.6%;
                                                                                                                     16 G; 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pred. No. 1.9e+02; 0; Mismatches 3;
                                                         0
                                                       Score 42.2; DB 1;
Pred. No. 1.9e+02;
0; Mismatches 3;
                                                                                                                     T; 0
                                                                                                                     U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       genetic testing
                                                                                    Length 51;
                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              713
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                                                       Gaps
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RESULT 98
ACC84458
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                           S
                                                             Matches
                                                                        Query Match
Best Local
                                                                                                                                              The present invention relates to a neural thread protein (NTP) peptide referred to as cell death peptide. Thought to be cytostatic, antibacterial, immunosuppressive and antiinflammatory. It is useful for treating a condition in a patient requiring removal or destruction of cells, for treating a condition such as benign or malignant tumor, inflammatory disease, autoimmune disease and infectious disease. The peptide useful for treatment is derived from the amino acid sequence for a pancreatic thread protein. The peptide is conjugated, linked or bound to a molecule chosen from antibody or its fragment, antibody-like binding molecule, where the molecule has a higher affinity for binding to a tumor or other target than binding to other cells. Treatment using NTP peptides can remove benign tumors with less risk and fewer of the undesirable side effects of surgery. The present sequence is an NTP encoding sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19-JUL-2001;
19-JUL-2001;
16-NOV-2001;
                                                                                                                                                                                                                                                                                                                                                                                                       Novel neural thread protein peptide, referred as cell death peptide, useful for treating prostatic hyperplasia, psoriasis, eczema, dermatosis, atherosclerosis, cosmetic modification to skin, throat, mouth, muscle.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NTP peptide
                                                                                                                       Sequence 42 BP; 8 A; 15 C; 10 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                          Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      P-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Averback PA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WC2003008443-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Cytostatic; Antibacterial; Immunosuppressive; Antiinflammatory; neural thread protein; NTP; tumour; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                28-AUG-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ACC84458;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ACC84458 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19-JUL-2002; 2002WO-CA001105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-JAN-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  neural thread protein; NTP; tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                XOMXN (-OMXN)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2003-247999/24.
                369 TCCACCTGCCTCAGCCTCCCAAAGTGCTGGGATTACAGGCGT 410
                                                           42;
μ
                                                                           4.2%;
Similarity 100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABR63253
TCCACCTGCCTCAGCCTCCCAAAGTGCTGGGATTACAGGCGT
                                                                                                                                                                                                                                                                                                                                                                        Page 16; 77pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; 2001US-0306150P.
; 2001US-0306161P.
; 2001US-0331477P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 encoding
                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sequence
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                                                                                                                                                                                                                                                                                                                                                                          English.
                                                           0,
                                                                           Score 42;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   #5
                                                             Mismatches
                                                                                            DB 1;
                                                                             1.6e+02;
                                                             0
                                                                                        Length 42;
                                                             Indels
                                                           0
                                                             Gaps
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28-AUG-2003 ACC84457; ACC84457

(first entry)

DNA;

42 BP

NTP

peptide encoding

sequence

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RESULT 100
ACC84459
ID ACC844
XX
ACC844

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Best Local S
Matches 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      referred to as cell death peptide. Thought to be cytostatic, antibacterial, immunosuppressive and antiinflammatory. It is useful for treating a condition in a patient requiring removal or destruction of cells, for treating a condition such as benign or malignant tumor, inflammatory disease, autoimmune disease and infectious disease. The peptide useful for treatment is derived from the amino acid sequence for a pancreatic thread protein. The peptide is conjugated, linked or bound to a molecule chosen from antibody or its fragment, antibody-like binding molecule, where the molecule has a higher affinity for binding to a tumor or other target than binding to other cells. Treatment using NTP peptides can remove benign tumors with less risk and fewer of the undesirable side
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel neural thread protein peptide, referred as cell death peptide, useful for treating prostatic hyperplasia, psoriasis, eczema, dermatos atherosclerosis, cosmetic modification to skin, throat, mouth, muscle.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 16; 77pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2003-247999/24.
P-PSDB; ABR63252.
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19-JUL-2001; 2001US-0306161P.
16-NOV-2001; 2001US-0331477P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    effects of surgery. The present sequence is an NTP encoding sequence
                                                                                                                                                                                                                                                                   NTP peptide encoding sequence
                                                                                                                                                                                                                                                                                                                              28-AUG-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                ACC84459 standard; DNA; 42
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                                                                WO2003008443-A2
                                                                                                                                                                                neural
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42; Conservative
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Pred. No.
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hes 0;
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20-OCT-2000; 2000US-0241994P. 08-JUN-2001; 2001US-0296764P. 22-OCT-2001; 2001WO-US047856

(BIOC-) BIOCARDIA INC

25-JUL-2002.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel neural thread protein peptide, referred as cell death peptide, useful for treating prostatic hyperplasia, psoriasis, eczema, dermat atherosclerosis, cosmetic modification to skin, throat, mouth, muscl
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19-JUL-2001; 2001US-0306161P.
16-NOV-2001; 2001US-0331477P.
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                                                                                                                                                                                                                                                   T7; leukocyte; gene expression profiling; allograft rejection; atherosclerosis; congestive heart failure; systemic lupus erytrheumatoid arthritis; osteoarthritis; cytomegalovirus; infecti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to a neural thread protein (NTP) peptide referred to as cell death peptide. Thought to be cytostatic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 17; 77pp; English.
                                                                                                                                                                                                                                                                                                                                                            09-JAN-2003
                                                                                                                                                                                                                                                                                                                                                                                          ABZ07763;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 42 BP; 6 A; 16 C; 11 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      P-PSDB; ABR63254.
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                                                                                                                                                                                                                                                                                                                        Human leukocyte gene expression profiling probe
                                                                                                                                                                                                                                                                                                                                                                                                                             ABZ07763 standard; DNA; 50
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1;
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                                                                                                                                                                                                                                                                                                                              ŏ
                                                                                                                                                                                                                                                          infection; probe;
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Best Local
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                           The
                                                                                                                                                                                                                                                                                                                                                                                  gene
mRNAs encoding sed
identified within
                                                      Claim 1; SEQ ID NO 18997; 71pp + Sequence Listing; English.
                                                                                New nucleic acid that is a 5' expressed sequence tag (5' EST) fobtaining cDNAs and genomic DNAs that correspond to 5'ESTs and diagnostic, forensic, gene therapy and chromosome mapping proce
                                                                                                                                               WPI;
                                                                                                                                                                         Dumas Milne Edwards J,
                                                                                                                                                                                                                                  26-FEB-1999;
                                                                                                                                                                                                                                                               21-FEB-2000; 2000EP-00200610.
                                                                                                                                                                                                                                                                                            06-SEP-2000.
                                                                                                                                                                                                                                                                                                                         EP1033401-A2
                                                                                                                                                                                                                                                                                                                                                      Homo
                                                                                                                                                                                                                                                                                                                                                                                                 Human; 5'
                                                                                                                                                                                                                                                                                                                                                                                                                              Human secreted
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAC14922 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New system for leukocyte expression profiling, diagnosing a disease, or monitoring (the rate of) progression of a disease, e.g. atherosclerosis or congestive heart failure, comprises diagnostic oligonucleotides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Wohlgemuth J, I
             present sequence is one of a large number of 5' ESTs derived as encoding secreted proteins. No ORF has yet been conclusivel
                                                                                                                                                                                                                                                                                                                                                                                n; 5' EST; expressed sequence tag; secreted
therapy; chromosome mapping; ss.
                                                                                                                                               2000-500381/45
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                                                                                                                                                                                                                                                                                                                                                                                                                            protein 5' EST,
                                                                                                                                                                                                                                    99US-0122487P
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Pred. No. 1.9e
0; Mismatches
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                                                                                                                                                                          Giordano
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            been conclusively
                                                                                  mapping procedures
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This invention relates to novel isolated nucleotide sequences which comprise 217 defined polymorphic sequences. Sequence polymorphism-based analysis of nucleic acid sequences can augment or replace previously known methods for determining the identity and relatedness of

Claim

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sequence polymorphism analysis; human identity; human relatedness; single nucleotide polymorphism; SNP; genetic disease; cytostatic; immunosuppressive; antiniflammatory; neuroprotective; antinicrobial; fatty acid metabolism; glycolysis; amino acid metabolism; glycolysis; amino acid metabolism; patternity analysis; forensic; autoimmune disease; cancer; nervous sysinfection; pathogenic microorganism; human; ds.
                                                                                                                                                                                                                                          Novel polynucleotide and polypeptide including one or more substitute including one or more substituted including and treating associated with the presence of sequence polymorphism in humbers.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17-NOV-1999;
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99US-00443199
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/*tag= a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
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90.0%;
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Pred. No. 1.9e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
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RESULT 104
AAL27794/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cc individuals. Single nucleotide polymorphisms (SNPs) tend to occur with cc great frequency throughout the genome and may be located close to loci of cc interest. Such variations can cause or be closely linked to pathological conditions (genetic diseases). Hence the SNPs of the invention may be considually in the development of compounds with cytostatic, antimicrobial continuosuppressive, antiniflammatory, neuroprotective or antimicrobial continuosuppressive, antiniflammatory, neuroprotective or antimicrobial continuosuppressive, antiniflammatory, neuroprotective or antimicrobial continuosuppressive, and amino acid metabolism may also be developed. The compounds may be useful for treating a subject suffering from or at crisk for a pathology associated with the presence of a sequence colymorphism. SNP detection is also useful in paternity analysis and corganism and phenotypic traits include genetic diseases such as corganism and phenotypic traits include genetic diseases such as corganism and infection by pathogenic microorganisms. The present sequence is the sequence corrowed and including a human SNP of the invention.
Query Match
Best Local S
Matches 45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Immunosuppressive; immunostimulatory; antiinflammatory; cytostatic; neuroprotective; antimicrobial; gene therapy; vaccine; amylase; cancer; amyloid protein; angiopoietin; apoptosis related protein; cadherin; cyclin; polymerase; oncogene; histone; kinase; colony stimulating factor; complement related protein; cytochrome; kinesin; cytokine; interferon; interleukin; G-protein coupled receptor; thioesterase; inflammation; multifactorial disease; autoimmune disease; infection; nervous system disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 51 BP; 12 A; 11 C; 17 G; 11 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAL27794 standard; DNA; 51
                                                                                                                                                                                                                                                                                 28-DEC-1999; 99US-0173419P.
27-DEC-2000; 2000US-00173419.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human SNP
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                                                                                                                                                                            WPI; 2001-465210/50
                                                                                                                                                                                                             Shimkets
                                                                                                                                                                                                                                                                                                                                     28-DEC-2000; 2000WO-US035498
                                                                                                                                                                                                                                                                                                                                                                                                           WO200147944-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                (CURA-) CURAGEN CORP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TGAGCTCAAGCAGTCCACCTGCCTCAGCCTCCCAAAGTGCTGGGATTACA 405
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                                                                                                                                                                                                             Leach M;
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90.0%;
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Pred. No.
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The present invention relates to oligonucleotides variants of proteins related to amylases, amyloid apoptosis related proteins, cadherin, cyclin, poly

ides encoding polymorphic loid proteins, angiopoietin, polymerase, oncogenes,

Claim 1;

Page 1666; 4143pp; English.

Polymorphic nucleic acids encoding e.g. amylases, cyclins, polymerases, oncogenes and histones, useful for diagnosing and treating, e.g. cancer, autoimmune diseases and infections.

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ARESULT 105
AAA173932
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protein coupled receptors and thioesterases. The present sequence is one
such oligonuclectide. The oligonuclectides and the peptides encoded by
them may be used in the prevention, diagnosis and treatment of diseases
associated with inappropriate expression of the proteins listed above.
Disorders that may be prevented, diagnosed and/or treated include
multifactorial diseases with a genetic component, such as autoimmune
diseases (e.g. rheumatoid arthritis, multiple sclerosis, diabetes,
systemic lupus erythromatosus and Grave's disease), inflammation, cancer
(e.g. cancers of the bladder, brain, breast, colon and kidney,
leukaemia), diseases of the nervous system and an infection of pathogenic
AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide sequences (I), which contain single nucleotide polymorphisms (SNPs). AAM53114 to AAM5329 represent peptides related to human polymorphic polymorphic polymorphic sequences. The sequences can be used in gene and protein therapy, and in vaccine production. (I) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of polymorphic polypeptides. For example, (I) may be used to treat disorders by rectifying mutations or deletions in a patient's genome that affect the activity of polypeptides by expressing inactive proteins or to supplement the patients own production of polypeptide. Additionally, (I) and its complementary sequences may also be used as DNA probes in diagnostic assays to detect
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      histones, kinases, colony stimulating factors, complement proteins, cytochromes, kinesins, cytokines, interferons, i
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29-NOV-2000; 2000US-00726173.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; single nucleotide polymorphism;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human silent SNP containing nucleic acid SEQ:873
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAI73932 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                Claim 1; Page 321; 2653pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Polymorphic nucleic acid sequences, useful in genetic testing and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2001-356160/37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Shimkets RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (CURA-) CURAGEN CORP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Leach M;
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restorative therapy; polymorphic;
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Pred.
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RESULT 106
                        CC AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide CC sequences (I), which contain single nucleotide polymorphisms (SNPs). CC AAM53114 to AAM53329 represent peptides related to human polymorphic CC thamsall14 to AAM53329 represent peptides related to human polymorphic CC polynucleotide sequences. The sequences can be used in gene and protein CC therapy, and in vaccine production. (I) and the polypeptides encoded by CC them may be used in the prevention, diagnosis and treatment of diseases CC associated with inappropriate expression of polymorphic polypeptides. For CC example, (I) may be used to treat disorders by rectifying mutations or CC example, (I) may be used to treat disorders by rectifying mutations or CC polymersing inactive proteins or to supplement the patients own CC production of polypeptide. Additionally, (I) and its complementary CC sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and CC therefore which patients may be in need of restorative therapy. The CC polypeptides encoded by (I) may be used as antigens in the production of antibodies may also be used to down regulate expression and activity. The antibodies may also be used to down regulate expression and activity. The antibodies may also
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Polymorphic nucleic acid sequences, useful in genetic testing
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29-NOV-2000;
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as diagnostic agents for detecting the presence of
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2000US-00726173.
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90.0%;
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0; Mismatches
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Query Match Best Local Similarity

4.2%;

Score 42; Pred. No.

DB 1; 1.9e+02;

Length 51;

Sequence

51 BP;

18 A; 13 C;

10 G; 10 T; 0 U; 0 Other;

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RESULT 107
AAI77521/c
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                           CC AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide CC sequences (I), which contain single nucleotide polymorphisms (SNPs). CC AAM53114 to AAM53129 represent peptides related to human polymorphic CC polynucleotide sequences. The sequences can be used in gene and protein therapy, and in vaccine production. (I) and the polypeptides encoded by CC them may be used in the prevention, diagnosis and treatment of diseases CC associated with inappropriate expression of polymorphic polypeptides. For CC example, (I) may be used to treat disorders by rectifying mutations or CC deletions in a patient's genome that affect the activity of polypeptides by expressing inactive proteins or to supplement the patients own CC production of polypeptide. Additionally, (I) and its complementary CC sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and CC polypeptides encoded by (I) may be used as antigens in the production of antibodies specific for polymorphic polypeptides. The antibodies may also be used to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic colypeptides.
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Best Local
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29-NOV-2000; 2000US-00726173
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les 45; Conserv
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Pred. No. 1.9e+02;
0; Mismatches 5;
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RESULT 108
RASIULY 108
ANAT79584/c
ID ANAT795
XX ANAT795
XX HAMAN
DT 09-NOV
XX HUMAN
KW PROTES
KW GUANTES
KW MPI;
KW GUANTES
CC GAMM53
CC GAMM54
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CC AAM533114 to AAM53329 represent peptides related to human polymorphic polynucleotide sequences. The sequences can be used in gene and protein therapy, and in vaccine production. (I) and the polypeptides encoded by thermapy be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of polymorphic polypeptides. For example, (I) may be used to treat disorders by rectifying mutations or deletions in a patient's genome that affect the activity of polypeptides by expressing inactive proteins or to supplement the patients own production of polypeptide. Additionally, (I) and its complementary sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and therefore which patients may be in need of restorative therapy. The polypeptides encoded by (I) may be used as antigens in the production of polypeptides specific for polymorphic polypeptides. The antibodies may also be used to down regulate expression and activity. The antibodies may also be used to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic
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Best Local :
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29-NOV-2000; 2000US-00726173
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 1; Page 2503; 2653pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Polymorphic nucleic acid sequences, useful in genetic testing
                                                                                                                                                                                                                                                     Sequence 51 BP; 10 A; 14 C; 18 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (CURA-) CURAGEN CORP.
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51
                                                                                                                            45;
                                                                                                                                                          Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TTTAGTAGAGATGGAGTTTCTCCATGTTGGTCAGGCTGGTCTCGAACTCC
                                                         AGGCTGGAGTGCAGTGGCGAATCTTGGCTCACTGCAACCTCTGCCTCCC 695
   AGGCTGGAGTGCAGTGGCGTGATCTCGGCTCACTGCAACCTCCACCTCCC 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
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                                                                                                                            Conservative
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                                                                                                                                                                                                                                                                                                               samples
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                                                                                                                                                    4.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
                                                                                                                            0,
                                                                                                                     Score 42; DB 1; Le
Pred. No. 1.9e+02;
0; Mismatches 5;
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                                                                                                                                                                                         Length 51;
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RESULT 109
AAI78039/c
AAI78039/c
XX
AAI78039/s
AC AAI78039;
AC CLaim 1;
AC AAI73060;
AC AAI73060
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                                                                                                                                                                                                                                                                                                                                                         AAI73060 to AAI79867 represent isolated human polymorphic polymucleotide sequences (I), which contain single nucleotide polymorphisms (SNPs). CC AAM53114 to AAM5329 represent peptides related to human polymorphic polymorphic golymucleotide sequences. The sequences can be used in gene and protein therapy, and in vaccine production. (I) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases CC associated with inappropriate expression of polymorphic polypeptides. For CC deletions in a patient's genome that affect the activity of polypeptides or CC by expressing inactive proteins or to supplement the patients own CC production of polypeptide. Additionally, (I) and its complementary CC sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and CC therefore which patients may be used as antigens in the production of polypeptides encoded by (I) may be used as antigens in the production of CC antibodies specific for polymorphic polypeptides. The antibodies may also be used to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic constraints of polymorphic polypeptides.
                                                                                                                                       Matches
                                                                                                                                                                       Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; single nucleotide polymorphism; SNP; genome; gene ther protein therapy; vaccine; probe; diagnostic assay; detection; quantitation; restorative therapy; polymorphic; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  09-NOV-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-356160/37.
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29-NOV-2000; 2000US-00726173.
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                                                                                                                                                                                                                                                                    Sequence 51 BP; 10 A; 13 C; 17 G; 11 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; Page 2034; 2653pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Polymorphic nucleic acid sequences, useful in genetic testing
                                                                                                                                                                                                                                                                                                                                           polypeptides in samples
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (CURA-) CURAGEN CORP.
                                          843 CCTGCCTCGGCCTCCCAAAGTGCTGGGATTACAGGCGTGAGCCACCACGC 892
       50
                                                                                                                                       45;
                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          standard; DNA; 51 BP
CCTGCCTCAGCCTCCCAAAGTGCTGGGATTACAGGCATGAGCCACTGTGC
                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
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                                                                                                                                                                       90.0%;
                                                                                                                                       Score 42; DB
Pred. No. 1.9e
0; Mismatches
                                                                                                                                           0
                                                                                                                                                                                                              DB 1;
                                                                                                                                                                                  9e+02;
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                                                                                                                                                                                                       Length 51
                                                                                                                                              Indels
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RESULT 110 AAI78300 ID AAI783 XX AC AAI783

AAI78300 standard; DNA; 51

AA178300

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EXEXEXEX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 111
AAI73860
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CC sequences (I), which contain single nucleotide polymorphisms (SNPs).

CC AAW53114 to AAW5329 represent peptides related to human polymorphic polynucleotide sequences. The sequences can be used in gene and protein therapy, and in vaccine production. (I) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases consisted with inappropriate expression of polymorphic polypeptides. For cexample, (I) may be used to treat disorders by rectifying mutations or converse in a patient's genome that affect the activity of polypeptides by expressing inactive proteins or to supplement the patients own converse in a patient's genome that affect the activity of polypeptides by expressing inactive proteins or to supplement the patients own converse may also be used as NNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and converse which patients may be in need of restorative therapy. The polypeptides encoded by (I) may be used as antigens in the production of antibodies may also be used as diagnostic agents for detecting the presence of polymorphic converse and activity. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic converse in amounter agents for detecting the presence of polymorphic converse in amounter agents for detecting the presence of polymorphic converse in amounter agents for detecting the presence of polymorphic converse in amounter agents for detecting the presence of polymorphic converse in amounter agents for detecting the presence of polymorphic converse in amounter agents for detecting the presence of polymorphic converse in a paralles.
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                  Matches
Human; single nucleotide polymorphism; SNP; genome; gene therapy;
                                       Human silent SNP containing nucleic acid SEQ:801.
                                                                                      09-NOV-2001
                                                                                                                                                                        AAI73860 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; Page 2114; 2653pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Polymorphic nucleic acid sequences, useful in genetic testing and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2001-356160/37
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                30-NOV-1999; 99US-0168138P.
29-NOV-2000; 2000US-00726173.
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                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
                                                                                                                                                                                                                                                                                                                      847 CCTCGGCCTCCCAAAGTGCTGGGATTACAGGCGTGAGCCACCACGCCCGG 896
                                                                                                                                                                                                                                                                                                                                                                                                                                                       5
                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP; 12
                                                                                                                                                                                                                                                                                                                                                                  Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SNP containing nucleic acid SEQ:5241.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 samples
                                                                                                                                                                        DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; 17 C; 14 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                      4.2%;
90.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           represent isolated human polymorphic polynucleotide contain single nucleotide polymorphisms (SNPs).
                                                                                                                                                                          51
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                                                                                                                                                                                                                                                                                                                                                                                      Score 42; DB 1;
Pred. No. 1.9e+02
                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                    1.9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                          Length 51;
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                                                                                                                                                                                                                                                                                51
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Homo sapiens. WO200140521-A2

Human; single nucleotide polymorphism; SNP; genome; gene therapy; protein therapy; vaccine; probe; diagnostic assay; detection; quantitation; restorative therapy; polymorphic; ds.

Human silent SNP containing nucleic acid SEQ:701.

09-NOV-2001

(first entry)

AAI73760 standard; DNA; 51

ВP

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Query Match Best Local : Matches 4:

45;

Conservative

Similarity

4.2%;

Score 42; DB 1; Leng; Pred. No. 1.9e+02;

Length 51, ; 5; Indels

<u>,</u>

Gaps

0

638 TGTCACCCAGGCTGGAGTGCAGTGGCGCAATCTTGGCTCACTGCAACCTC

TGTCACCCAGGCTGAACTGCAGTGGTGATCTTGGCTCACTGCAACCTC

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CC AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide
CC sequences (I), which contain single nucleotide polymorphisms (SNPs).
CC AAM53114 to AAM53329 represent peptides related to human polymorphic
CC polynucleotide sequences. The sequences can be used in gene and protein
CC therapy, and in vaccine production. (I) and the polypeptides encoded by
CC them may be used in the prevention, diagnosis and treatment of diseases
CC associated with inappropriate expression of polymorphic polypeptides. For
CC deletions in a patient's genome that affect the activity of polypeptides
CC to example, (I) may be used to treat disorders by rectifying mutations or
CC deletions in a patient's genome that affect the patients own
CC production of polypeptide. Additionally, (I) and its complementary
CC sequences may also be used as DNA probes in diagnostic assays to detect
CC antibodies specific for polymorphic polypeptides. The
CC therefore which patients may be used as antigens in the production of
CC polypeptides encoded by (I) may be used as antigens in the production of
CC be used to down regulate expression and activity. The antibodies may also
De used as diagnostic agents for detecting the presence of polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-NOV-1999; 99US-0168138P.
29-NOV-2000; 2000US-00726173.
Sequence 51 BP; 9 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2001-356160/37
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quantitation; restorative therapy; polymorphic; ds.
                                        polypeptides in samples
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Polymorphic nucleic acid sequences, useful in genetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3
16 C; 13 G; 13 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             testing
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide sequences (I), which contain single nucleotide polymorphisms (SNPs). CC AAM53114 to AAM53129 represent peptides related to human polymorphic polynucleotide sequences. The sequences can be used in gene and protein therapy, and in vaccine production. (I) and the polypeptides encoded by CC them may be used in the prevention, diagnosis and treatment of diseases CC associated with inappropriate expression of polymorphic polypeptides. For CC deletions in a patient's genome that affect the activity of polypeptides or CC deletions in a patient's genome that affect the patients own CC by expressing inactive proteins or to supplement the patients own CC sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and CC therefore which patients may be in need of restorative therapy. The CC antibodies may be considered as antigens in the production of CC antibodies specific for polymorphic polypeptides. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic polypeptides. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic.
                                                                                                                                                                                                                                                                                                                                                 AAI77806
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29-NOV-2000; 2000US-00726173
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2001-356160/37
                                                                                                                                                                Human; single nucleotide polymorphism; SNP; genome; protein therapy; vaccine; probe; diagnostic assay; quantitation; restorative therapy; polymorphic; ds.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Polymorphic nucleic acid sequences, useful in genetic testing
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                                                                                                                                                                                                                                                                 09-NOV-2001
                                                                                                                                                                                                                                                                                                 AAI77806
                                                                                                                                                                                                                                                                                                                                  AAI77806 standard; DNA; 51
 30-NOV-1999;
                               30-NOV-2000; 2000WO-US032758
                                                                  07-JUN-2001
                                                                                                 WO200140521-A2
                                                                                                                                                                                                                              Human silent SNP containing nucleic acid SEQ:4747.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      polypeptides in samples
                                                                                                                                                                                                                                                                                                                                                                                                                                                      648
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                                                                                                                                                                                                                                                                                                                                                                                                                                          GCTGGAGTGCAGTGGCGCAATCTTGGCTCACTGCAACCTCTGCCTCCCGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4.2%;
larity 90.0%;
Conservative
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 99US-0168138P
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Pred. No. 1.9e+02;
0; Mismatches 5
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                                                                                                                                                                                 detection
                                                                                                                                                                                                 gene therapy;
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RESULT 114
AAI73533/c
ID AAI735
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AAI73533/c
ID AAI735
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DT 09-NOV
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Human
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Human
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; single nucleotide polymorphism; SNP; genome; gene ther protein therapy; vaccine; probe; diagnostic assay; detection; quantitation; restorative therapy; polymorphic; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 51 BP; 7 A; 22 C; 9 G; 13 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; Page 1963; 2653pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-356160/37.
                                                                                                                                                                30-NOV-1999; 99US-0168138P
29-NOV-2000; 2000US-00726173
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                                                                                                                                                                                                                                                                                                                             07-JUN-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       672 GGCTCACTGCAACCTCTGCCTCCCGGGTTCAAGTTATTCTCCTGCCCCAG
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Pred. No. 1.9e
0; Mismatches
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Shimkets RA,

Leach M

(CURA-) CURAGEN CORP.

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CC AAM53114 to AAM5329 represent peptides related to human polymorphic in abmples, and continuous particles encoded by (I) may be used as an attibodies may also be used to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic polymorphic polymorphic.
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Best Local S
Matches 45
Isolated human nucleic acids comprising one or more single nucleotide polymorphisms, useful for treating a subject suffering from a pathology, e.g. autoimmune diseases, ascribed to the presence of a sequence polymorphism.
                                                                                                                                                                              24-NOV-1999;
                                                                                                                                                                                                                                            31-MAY-2001
                                                                                                                                                                                                                                                                            WO200138586-A2
                                                                                                                                                                                                                                                                                                                                                      Human; single nucleotide polymorphism; SNP; polymorphism; cytostatic; immunosuppressive; antiinflammatory; neuroprotective; antimicrobial; autoimmune disease; inflammation; cancer; nervous system disease;
                                                                                                                                                                                                                                                                                                                                                                                                                       Human silent
                                                                                                                                                                                                                                                                                                                                                                                                                                                       05-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABL00195 standard; DNA; 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1; Page 199; 2653pp; English
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                                                                                                                                                                                                           22-NOV-2000; 2000WO-US032311
                                                                                                                                                                                                                                                                                                                                       autoimmune disease; inflammation; cinfection; polymorphic protein; ds.
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45; Conser
                                                                                                                                              CURAGEN CORP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GCCCGCCTCGGCCTCCCAAAGTGCCGGGATTACAGGCTTGAGTCACCATG
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                                                                                                                                                                                                                                                                                                                                                                                                                       noncoding
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                                                                                                              Leach
                                                                                                                                                                            99US-0167383P
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                                                                                                                                                                                                                                                                                                                                                                                                                       SNP oligonucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 42;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
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1.9e+02;
5;
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                                                                                                                                                                                                                                                                                                                                                                                                                       SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 51;
                                                                                                                                                                                                                                                                                                                                                                                                                         NO:186
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ABLO0010 to ABL01104 represent human nucleic acid oligonucleotides comprising one or more single nucleotide polymorphisms (SNPs). ABB56531 to ABB56903 represent human peptides encoded by some of the SNP oligonucleotides. The sequences from the present invention can have immunosuppressive, cytostatic, antiinflammatory, neuroprotective and antimicrobial activities. Nucleic acids, polypeptides, oligonucleotides and antibodies from the present invention can be used for treating a subject suffering from, at risk for, or suspected of, suffering from a pathology may be autoimmune diseases, inflammation, cancer, diseases of the nervous system, and infection by pathogenic microorganisms. The SNPs are also useful for determining which forms of a characterised in polymorphism are present in individuals. The antibodies may be used in
                                                                                  the detection, quantitation and/or cellular or tissue localisation of a golymorphic protein (e.g., for use in measuring levels of the polymorphic protein within appropriate physiological samples)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1;
        Sequence 51
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        BP; 8
        P.
        23 C;
        œ
G;
        12 T; 0 U; 0 Other;
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RESULT 116
ADDI12525/c
ID ADDI125
XX ADDI125
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Best Local S
Matches 45
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breast cancer susceptibility gene 1; BRCAl; repetitive Alu;
ovarian cancer; junction sequence; recombination; mutant.
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45; Conserv
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llarity 90.0%;
Conservative
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Pred. No. 1.9e
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1.9e+02;
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This invention relates to a novel method for predicting a predisposition to cancer in a patient by detecting large deletions in the human tumour suppressor gene identified as the breast cancer susceptibility gene 1 (BRCA1). Specifically, it refers to deletions that result from the unequal crossover between a pair of repetitive Alu sequences in the BRCA.

the BRCAI

Predicting a predisposition to cancer in a patient comprising detecting deletion in the BRCA1 gene that results from the unequal crossover between a pair of repetitive sequences in the BRCA1 gene.

SEQ

ID NO

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59pp; English.

WPI; 2004-062369/06

07-JUN-2002; 09-AUG-2002;

2002US-0387132P 2002US-0402430P

(MYRI-)

MYRIAD GENETICS INC.

Hendrickson

BC,

Ward

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Pruss

09-JUN-2003; 2003WO-US018098

WO2003104474-A2

18-DEC-2003

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RESULT 117
AAI75849
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Best Local
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AAIT3060 to AAIT9867 represent isolated human polymorphic polynucleotide sequences (I), which contain single nucleotide polymorphisms (SNPs). AAM53114 to AAM5329 represent peptides related to human polymorphic polynucleotide sequences. The sequences can be used in gene and protein therapy, and in vaccine production. (I) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of polymorphic polypeptides. For deletions in a patient's genome that affect the activity of polypeptides by expressing inactive proteins or to supplement the patients own production of polypeptide. Additionally, (I) and its complementary sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and therefore which patients may be in need of restorative therapy. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-NOV-1999; 99US-0168138P
29-NOV-2000; 2000US-00726173
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; single nucleotide polymorphism; SNP; genome; gene ther protein therapy; vaccine; probe; diagnostic assay; detection; quantitation; restorative therapy; polymorphic; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human silent SNP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             09-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAI75849 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     and 16, given in an exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2001-356160/37.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30-NOV-2000; 2000WO-US032758
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Polymorphic nucleic acid sequences, useful in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (CURA-) CURAGEN CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        696
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GGGTTCAAGCAATTCTCCTGCCTCAGCCTCCTGAGTAGCTGGGATTAC 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Leach
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2653pp; English
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Pred. No. 1
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                                                                                  comprising one or more single nucleotide polymorphisms (SNPs). ABB56531 to ABB56903 represent human peptides encoded by some of the SNP coligonucleotides. The sequences from the present invention can have antimicrobial activities. Nucleic acids, polypeptides, oligonucleotides and antibodies from the present invention can be used for treating a subject suffering from, at risk for, or suspected of, suffering from a pathology ascribed to the presence of a sequence polymorphism. The pathology may be autoimmune diseases, inflammation, cancer, diseases of the nervous system, and infection by pathogenic microorganisms. The SNPs care also useful for determining which forms of a characterised colymorphism are present in individuals. The antibodies may be used in the detection, quantitation and/or cellular or tissue localisation of a concerning from the conjugation of a characterised colymorphic protein (e.g., for use in measuring levels of the polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human silent noncoding SNP oligonucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 polypeptides in samples
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Isolated human nucleic acids comprising one or more single nucleotide polymorphisms, useful for treating a subject suffering from a pathology, e.g. autoimmune diseases, ascribed to the presence of a sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            31-MAY-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             autoimmune disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; single nucleotide polymorphism; SNP; polymorphism; cytostatic; immunosuppressive; antiinflammatory; neuroprotective; antimicrobial;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 05-MAR-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; Page 286; 674pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       24-NOV-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          infection; polymorphic protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABL00010 to ABL01104 represent human nucleic acid oligonucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              polymorphism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22-NOV-2000; 2000WO-US032311.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    966
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                                                                        within
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                                                                     appropriate physiological samples)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4.2%;
91.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  English
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Pred. No. 26
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                                                                                                                CC which contain single nucleotide polymorphisms (SNPs). Sequences 1112 (C (AAA76318 A77429) are consecutive pairs of nucleotides which contain C silent SNPs. Sequences 1113 to 1192 (AAA77430-A77509) are consecutive pairs of nucleotides containing SNPs which result in changes in the CC pairs of nucleotides containing SNPs which result in changes in the CC corresponding amino acid sequences (AAB11749-B11828). The SNPs in CC sequences 1113 to 1128 (AAA77430-A77445) lead to conservative amino acid CC changes, while those in sequences 1129 to 1186 (AAA77446-A77503) result in non-conservative changes. The SNPs in sequences 1187 to 1192 (AAA77504-A77509) generate frameshift mutations. The invention also CC relates to a method of detecting a polymorphic site in a nucleic acid and CC against such peptides containing polymorphic sites, antibodies raised CC groteins/peptides, and a method of detecting polymorphic CC gene therapy of an individual having, suspected of having, or at risk of CC developing a pathological condition due to the presence of a sequence CC polymorphism. Such treatment would comprise administration of the wild-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
Matches 44; Conser
         Sequence 51 BP; 8 A; 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequences AAA76318-A77509 represent 1192 human nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1; Page 498; 543pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         presence of a sequence polymorphism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human nucleic acids containing single nucleotide polymorphisms, us
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2000-387826/33.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Shimkets RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17-NOV-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human Alusubfamily SQ gene polymorphic site, SEQ ID NO:1125
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  51
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                                                                acid sequence. Antibodies raised against polymorphic also be used in the treatment of such individuals
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99US-00443199.
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         C; 15 G;
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Pred. No. 2e+02;
         20 T;
         0 U; 0 Other;
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RESULT 120
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Best Local (
                                                     changes, while those in sequences 1129 to 1186 (AAA77446-A77503) result in non- conservative changes. The SNPs in sequences 1187 to 1192 (AAA77504-A77509) generate frameshift mutations. The invention also relates to a method of detecting a polymorphic site in a nucleic acid and a method of detecting polymorphic sites, antibodies raised against such peptides containing polymorphic sites, antibodies raised against such peptides, and a method of detecting polymorphic proteins/peptides using the antibodies. The mucleic acids are useful for gene therapy of an individual having, suspected of having, or at risk of developing a pathological condition due to the presence of a sequence polymorphism. Such treatment would comprise administration of the wildtype nucleic acid sequence. Antibodies raised against polymorphic peptides can also be used in the treatment of such individuals
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequences AAA76318-A77509 represent 1192 human nucleic acid sequences which contain single nucleotide polymorphisms (SNPs). Sequences 1 to 1112 (AAA76318-A77429) are consecutive pairs of nucleotides which contain silent SNPs. Sequences 1113 to 1192 (AAA77430-A77509) are consecutive pairs of nucleotides containing SNPs which result in changes in the corresponding amino acid sequences (AABA1749-B11828). The SNPs in sequences 1113 to 1128 (AAA77430-A77445) lead to conservative amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human nucleic acids containing single nucleotide polymorphisms, us for treating a subject suffering, or at risk from a pathology due presence of a sequence polymorphism.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 1; Page 360; 543pp; English.
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99US-00443199.
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useful in the development of compounds with cytostatic, useful in the development of compounds with cytostatic, immunosuppressive, antiinflammatory, neuroprotective or antimicrobial activities. Regulators of metabolic pathways such as fatty acid metabolism, glycolysis, and amino acid metabolism may also be developed. The compounds may be useful for treating a subject suffering from or at risk for a pathology associated with the presence of a sequence polymorphism. SNP detection is also useful in paternity analysis and forensic application. Polymorphisms may contribute to the phenotype of an organism and phenotypic traits include genetic diseases such as autoimmune diseases, cancer, diseases of the nervous system and infection

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Best Local Similarity
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                                                                                                                                                                                                                                                                                 Human clone cg43089031 polymorphic site, SEQ ID NO:704.
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Pred. No. 2e+02;
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17-NOV-1998; 16-NOV-1999; 17-NOV-1999; 25-MAY-2000 WO200029623-A2 Shimkets RA, (CURA-) CURAGEN CORP. 2000-387826/33 Leach 98US-0109024P 99US-00443199 99WO-US027293

Human nucleic acids containing single nucleotide polymorphisms, useful for treating a subject suffering, or at risk from a pathology due to the presence of a sequence polymorphism.

Claim 1; Page 370; 543pp; English.

Sequences AAA76318-A77509 represent 1192 human nucleic acid sequences which contain single nucleotide polymorphisms (SNPs). Sequences 1 to 1112 (AAA76318-A77429) are consecutive pairs of nucleotides which contain silent SNPs. Sequences 113 to 1192 (AAA77430-A77509) are consecutive pairs of nucleotides which contain corresponding amino acid sequences (AAB11749-B11828). The SNPs in sequences 113 to 1128 (AAA77430-A77445) lead to conservative amino acid changes, while those in sequences (AAB11749-B11828). The SNPs in sequences 113 to 1128 (AAA77430-A77445) lead to conservative amino acid changes, while those in sequences 1129 to 1186 (AAA77446-A77503) result in non- conservative changes. The SNPs in sequences 1187 to 1192 (AAA77504-A77509) generate frameshift mutations. The invention also relates to a method of detecting a polymorphic site in a nucleic acid and a method of determining the relatedness of two nucleic acids. It also a method of determining the relatedness of two nucleic acids. It also encompasses peptides containing polymorphic sites, antibodies raised against such peptides, and a method of detecting polymorphic proteins/peptides using the antibodies. The nucleic acids are useful for gene therapy of an individual having, suspected of having, or at risk of developing a pathological condition due to the presence of a sequence polymorphism. Such treatment would comprise administration of the wild-Type nucleic acid sequence. Antibodies raised peptides can also be used in the treatment of against such ind individuals

Sequence 51 B₽; 12 A; 14 C; 16 G; 9 T; 0 U; 0 Other;

RESULT 122
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XX Homo s
XX FH Key
FT Variat 밁 Ś Best Local Similarity Matches 45; Conserv j. sequence polymorphism analysis; human identity; human single nucleotide polymorphism; SNP; genetic disease; immunosuppressive; antiinflammatory; neuroprotective; individuals. Single nucleotide polymorphisms (SNPs) tend to occur with great frequency throughout the genome and may be located close to loci of interest. Such variations can cause or be closely linked to pathological conditions (genetic diseases). Hence the SNPs of the invention may be This invention relates to novel isolated nucleotide sequences which comprise 217 defined polymorphic sequences. Sequence polymorphism-based analysis of nucleic acid sequences can augment or replace previously known methods for determining the identity and relatedness of Novel polynucleotide and polypeptide including one or more single nucleotide polymorphisms, useful for diagnosing and treating condit associated with the presence of sequence polymorphism in humans and Human single nucleotide polymorphism (SNP) region Seq ID32 18-DEC-2003 ADC16930; ADC16930 17-NOV-1999; WO200029622-A2 Homo sapiens fatty acid metabolism; glycolysis; amino acid metabolism;
paternity analysis; forensic; autoimmune disease; cancer; nervous system; immunosuppressive; fatty acid metaboli Claim 1; SEQ ID NO 32; 187pp; English animals. Shimkets RA, 17-NOV-1998; 16-NOV-1999; 25-MAY-2000 WPI; 2000-399731/34. variation infection; pathogenic microorganism; human; (CURA-) CURAGEN CORP 853 CCTCCCAAAGTGCTGGGATTACAGGCGTGAGCCACCACGCCCGGCTTATTT 903 51 standard; ccrccaaagracragaarraragacargaarcaccacaccraaccarri 1 Conservative (first entry Leach MD 98US-0109024P. 99US-00443199. 99WO-US027283 /standard_name= "Single nucleotide polymorphism" replace (26, T) Location/Qualifiers DNA; 88.2%; 51 ₽₽ 0, Pred. No. Mismatches 2e+02; 6 Indels cytostatic; antimicrobial; relatedness; 0; conditions Gaps

밁 S

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RASULT 123
AAIT6193/c
ID AAIT61
XX AAIT61
XX AAIT61
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XX PATCE
XX PATCE
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                                                                                                                                                                         CC polynucleotide sequences. The sequences can be used in gene and protein CC them may be used in the provention. (I) and the polypeptides encoded by CC them may be used in the prevention, diagnosis and treatment of diseases CC associated with inappropriate expression of polymorphic polypeptides. For CC example, (I) may be used to treat disorders by rectifying mutations or CC deletions in a patient's genome that affect the activity of polypeptides or production of polypeptide. Additionally, (I) and its complementary CC sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and therefore which patients may be in need of restorative therapy. The polypeptides specific for polymorphic polypeptides. The antibodies may also be used as diagnostic assents of complementary also be used to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic colypeptides. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic polymo
Query Match
Best Local Similarity
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Best Local
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                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide sequences (I), which contain single nucleotide polymorphisms (SNPs).

AAM53114 to AAM53329 represent peptides related to human polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; single nucleotide polymorphism; SNP; genome; gene ther protein therapy; vaccine; probe; diagnostic assay; detection; quantitation; restorative therapy; polymorphic; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 1; Page 1009; 2653pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Polymorphic nucleic acid sequences, useful in genetic testing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2001-356160/37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Shimkets RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30-NOV-1999; 99US-0168138P.
29-NOV-2000; 2000US-00726173.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30-NOV-2000; 2000WO-US032758.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human silent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    09-NOV-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         surrounding and including a human SNP of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (CURA-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1087 GAGGCGGGGTTTCACCATATTTGTCAGGCTGGTCTCAAACTCCTGACCTCA 1137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             4.2%;
1 Similarity 88.2%;
45; Conservation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURAGEN CORP.
                                                                                                                   51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GAGACGGGGTTTCACCATATTGGCCGGGATGGTCTCGAACTCCTGACCTCA 1
                                                                                                                   ₽,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SNP containing
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                                                                                                               13
                                                                                                               ð,
4.2%;
88.2%;
                                                                                                               16 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  51
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                                                                                                           13 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       nucleic acid SEQ:3134.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                14 G; 10 T; 0 U; 0 Other;
Score 41.4; DB 1; Length 51; Pred. No. 2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 41.4; DB 1;
Pred. No. 2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 6;
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RESULT 124
AAI73061/c
ID AAI730
XX AAI730
XX Human
XX Human
XX Human
XX Human
XX Human
XX Homo E
XX AO
PN WO2001
XX 30-NON
PR 29-NON
XX 30-NON
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XX (CURA-
XX WPI;
XX CLaim
XX 
                                                                                                                                                                                                                                                            CC sequences (I), which contain single nucleotide polymorphisms (SNPs).

CC AAM53114 to AAM53329 represent peptides related to human polymorphic polymorphic in the polymorphisms (SNPs).

CC polynucleotide sequences. The sequences can be used in gene and protein continuous the invaccine production. (I) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases consisted with inappropriate expression of polypeptides. For continuous in a patient's genome that affect the activity of polypeptides or conduction in a patient's genome that affect the activity of polypeptides by expressing inactive proteins or to supplement the patients own conduction of polypeptide. Additionally, (I) and its complementary considered as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and considered which patients may be used as antigens in the production of polypeptides encoded by (I) may be used as antigens in the production of bused to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic conclusions.
                                                                      Matches
                                                                                                  Query Match
Best Local
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                                                                                                                                                                                  Sequence 51
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                                                                                                                                                                                                                                            polypeptides in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; Page 54; 2653pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Polymorphic nucleic acid sequences, useful in genetic testing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2001-356160/37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Shimkets RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30-NOV-1999; 99US-0168138P.
29-NOV-2000; 2000US-00726173.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30-NOV-2000; 2000WO-US032758
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                                                                      45;
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                                                                                                  Similarity
CCTCTGCCTCCCGGGTTCAAGTTATTCTCCTGCCCCAGCCTCCTGAGTAGC 734
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TTAGTAGAGACGGGGTTTCACCATGCTGGCCAGGCTGGTCTCGAACTCCTG 1
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                                                                                                                                                                                     BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Leach M;
                                                                                                                                                                                     12
                                                                                                                                                                                     A; 11 C;
                                                                                            4.2%;
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                                                                   <u>,</u>
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                                                                                               Score 41.4;
Pred. No. 26
                                                                                                                                                                                     <u>ი</u>
                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                        σ
                                                                                                                                                                                     T; 0
                                                                                            2e+02;
                                                                                                                                                                                        U; 0 Other;
                                                                                                                           DB 1;
                                                                   6
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                                                                                                                           Length 51;
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                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; gene therapy;
detection;
                                                                   0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     and
                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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CCTCCGCCTCCTGGGTTCAAGCGATCCTCCTGCCTCAGCCTCCTGAGTAGC

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RESULT 126
AAI73736/c
ID AAI737
XX
AC AAI737
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                                                                                                                                                                                                                                                                            CC AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide CC sequences (I), which contain single nucleotide polymorphisms (SNPs). CC AAM53114 to AAM53129 represent peptides related to human polymorphic CC polynucleotide sequences. The sequences can be used in gene and protein therapy, and in vaccine production. (I) and the polypeptides encoded by CC them may be used in the prevention, diagnosis and treatment of diseases CC associated with inappropriate expression of polymorphic polypeptides. For CC deletions in a patient's genome that affect the activity of polypeptides or CC deletions in a patient's genome that affect the patients own CC by expressing inactive proteins or to supplement the patients own CC sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and CC antibodies specific for polymorphic polypeptides. The antibodies may also be used to down regulate expression and activity. The antibodies may also be used to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic polypeptides.
                                                                                                                                                                                                Query Match
Best Local S
                                                                                                                                                                                  Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Shimkets RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                30-NOV-1999; 99US-0168138P.
29-NOV-2000; 2000US-00726173.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human silent SNP containing nucleic acid SEQ:1716.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          09-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAI74775 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; Page 579; 2653pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2001-356160/37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30-NOV-2000; 2000WO-US032758
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200140521-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Polymorphic nucleic acid sequences, useful in genetic testing
   AAI73736;
                                                                                                                                                                                                                                                   Sequence
                                 AAI73736 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (CURA-) CURAGEN CORP.
                                                                                                                                                   997
                                                                                                                   L
                                                                                                                                                                                  45;
                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                     51
                                                                                                                                          GGCTCAAGCGATTCTCCTGTCTCAGCCTCCCAAGCAGCTGGGATTACGGGC 1047
                                                                                                                   GGCTCAAGCAATCCTCCCGCCTCAGCCTCCCAAGCAGCTGGGACTACAGGC 51
                                                                                                                                                                                                                                                   BP; 11
                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Leach M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first
                                                                                                                                                                                                4.2%;
                                                                                                                                                                                                                                                   A; 21 C; 12 G; 7 T; 0 U; 0 Other;
                                   51
                                                                                                                                                                                  0,
                                                                                                                                                                                  Score 41.4; DB 1; Length 51; Pred. No. 2e+02; O; Mismatches 6; Indels
                                                                                                                                                                                  0;
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                                                                                                                                                         RESULT 127
AAI76185/c
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                                                                                                                                                                                                                                                                                                                                                                                                    CC AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide CC sequences (I), which contain single nucleotide polymorphisms (SNPs). CC AAM33114 to AAM53329 represent peptides related to human polymorphic polynucleotide sequences. The sequences can be used in gene and protein therapy, and in vaccine production. (I) and the polypeptides encoded by CC them may be used in the prevention, diagnosis and treatment of diseases CC associated with inappropriate expression of polymorphic polypeptides. For CC example, (I) may be used to treat disorders by rectifying mutations or CC deletions in a patient's genome that affect the activity of polypeptides by expressing inactive proteins or to supplement the patients own CC production of polypeptide. Additionally, (I) and its complementary CC sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and CC polypeptides encoded by (I) may be used as antigens in the production of CC polypeptides specific for polymorphic polypeptides. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic be used as diagnostic agents for detecting the presence of polymorphic colypeptides may also be used as diagnostic agents for detecting the presence of polymorphic CC polypeptides in samples
                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                       Query Match
Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; single nucleotide polymorphism; SNP; genome; protein therapy; vaccine; probe; diagnostic assay; c quantitation; restorative therapy; polymorphic; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30-NOV-1999; 99US-0168138P: 29-NOV-2000; 2000US-00726173:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human silent SNP containing nucleic acid SEQ:677.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             09-NOV-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 1; Page 261; 2653pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Polymorphic nucleic acid sequences, useful in genetic testing and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2001-356160/37.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Shimkets RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          30-NOV-2000; 2000WO-US032758
Human; single nucleotide polymorphism; SNP; genome; gene therapy;
                                   Human silent
                                                                      09-NOV-2001
                                                                                                                                        AAI76185 standard; DNA; 51 BP
                                                                                                                                                                                                                                                                                                                                                                         Sequence 51 BP; 18 A; 11 C; 11 G; 11 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         therapy.
                                                                                                         AAI76185
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                    1052 GCCACCACACCCCGCTAATTTTTGTATTTCATTAGAGGCGGGGTTTCACC 1102
                                                                                                                                                                                                                                                                                                    45;
                                                                                                                                                                                                                                51
                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                GCCATCACCCCGGCTAATTTTTGTATTTTAGTAGAGACGGGGTTTCATC 1
                                                                                                                                                                                                                                                                                                    Conservative
                                   SNP containing nucleic acid SEQ:3126.
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                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                       4.2%;
                                                                                                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                                                                                                     Score 41.4; DB 1
pred. No. 2e+02;
                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                       DB 1;
                                                                                                                                                                                                                                                                                                      6
                                                                                                                                                                                                                                                                                                                                       Length 51;
                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           detection;
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                                                                                                                                                                                                                                                                                                        0,
                                                                                                                                                                                                                                                                                                          Gaps
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AAI74502/c
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             therapy, and in vaccine production. (1) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of polymorphic polypeptides. For example, (1) may be used to treat disorders by rectifying mutations or deletions in a patient's genome that affect the activity of polypeptides by expressing inactive proteins or to supplement the patients own production of polypeptide. Additionally, (1) and its complementary sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and therefore which patients may be in need of restorative therapy. The polypeptides encoded by (1) may be used as antigens in the production of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          protein therapy; vaccine; probe; diagnostic assay; detection;
quantitation; restorative therapy; polymorphic; ds.
WO200140521-A2
                                Homo sapiens
                                                                Human; single nucleotide polymorphism; SNP; genome; gene ther protein therapy; vaccine; probe; diagnostic assay; detection; quantitation; restorative therapy; polymorphic; ds.
                                                                                                                                 Human silent SNP containing nucleic acid SEQ:1443.
                                                                                                                                                                       09-NOV-2001
                                                                                                                                                                                                                                         AAI74502 standard; DNA; 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                polypeptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             antibodies specific for polymorphic polypeptides. The antibodies may also be used to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide sequences (I), which contain single nucleotide polymorphisms (SNPs). AAN53314 to AAN53329 represent peptides related to human polymorphic polynucleotide sequences. The sequences can be used in gene and protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1; Page 1007; 2653pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Polymorphic nucleic acid sequences, useful in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-356160/37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30-NOV-1999; 99US-0168138P.
29-NOV-2000; 2000US-00726173.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sapiens.
                                                                                                                                                                                                                                                                                                                                                              170
                                                                                                                                                                                                                                                                                                                              51
                                                                                                                                                                                                                                                                                                                                                                                            45;
                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                TTTTTTTTAGTAGAGATGGAGTTTCTCCATGTTGGTCAGGCTGGTCTCGA 220
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RΑ,
                                                                                                                                                                                                                                                                                                                              TGTATTTTTAGTAGAGACGGGGTTTCACCATGTTGGCCAGGCTGGTCTCGA 1
                                                                                                                                                                                                                                                                                                                                                                                           4.2%;
ilarity 88.2%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                in samples
                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Leach M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; 16 C;
                                                                                                                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                9 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             English
                                                                                                                                                                                                                                                                                                                                                                                                            Score 41.4; DB 1; Length 51; Pred. No. 2e+02;
                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                            6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               genetic testing
                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                 gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                              0
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RESULT 129
AAI76499
PRX PXX PXX SXX XXX XXX AXX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
                                                                                                                                                                           Human; single nucleotide polymorphism; SNP; genome; gene therapy; protein therapy; vaccine; probe; diagnostic assay; detection; quantitation; restorative therapy; polymorphic; ds.
                                                                                                                                                                                                                                                                                                                                                           AAI76499 standard; DNA; 51
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29-NOV-2000; 2000US-00726173
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                               30-NOV-2000; 2000WO-US032758
                                                                     07-JUN-2001.
                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                    Human silent
                                                                                                                                                                                                                                                                                      09-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 51 BP; 13 A; 16 C; 12 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 polypeptides in samples
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 1; Page 495; 2653pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Polymorphic nucleic acid sequences, useful in genetic testing and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2001-356160/37
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                                                                                                        WO200140521-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          178 TAGTAGAGATGGAGTTTCTCCCATGTTGGTCAGGCTGGTCTCGAACTCCCGA 228
                                                                                                                                                                                                                                                                                                                                                                                                                                                   51 TAGTAGAGATGGGGTTTTCGCCATGCTGGCCAGGCTGGTCTCAAACTCCTGA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          45;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RΑ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                      SNP
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                                                                                                                                                                                                                                                   containing nucleic acid SEQ:3440.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4.2%;
                                                                                                                                                                                                                                                                                                                                                             ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pred. No. 2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 41.4; DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                51
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30-NOV-1999;

99US-0168138P

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                                                                                                                                                                                                                                                                                                                                                                                                                            ID
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 130
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Best Local S
Matches 45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; Page 1103; 2653pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Polymorphic nucleic acid sequences, useful in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2001-356160/37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Shimkets RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29-NOV-2000; 2000US-00726173.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 51 BP; 10 A; 22 C; 11 G; 8 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                         Human; single nucleotide polymorphism; SNP; genome; gene ther protein therapy; vaccine; probe; diagnostic assay; detection; quantitation; restorative therapy; polymorphic; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          polypeptides in samples
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                                                                                                                                                                                                                                                                                                                            Human silent SNP
                                                                                                                                                                                                                                                                                                                                                             09-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                              AAI79633 standard; DNA; 51 BP
                                                                            30-NOV-1999; 99US-0168138P.
29-NOV-2000; 2000US-00726173.
                                                                                                                                                             07-JUN-2001.
                                                                                                                                                                                            WO200140521-A2
             Shimkets RA,
                                                                                                                           30-NOV-2000; 2000WO-US032758
                                              (CURA-) CURAGEN CORP.
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45; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CTCGAACTCCCGACCTCAGATGATCCCTCCGTCTCGGCCTCCCAAAGTGCT 266
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAI79867 represent isolated human polymorphic polynucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4.2%;
ilarity 88.2%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Leach M;
                                                                                                                                                                                                                                                                                                                                                             (first entry)
                 Leach M
                                                                                                                                                                                                                                                                                                                         containing nucleic acid SEQ:6574.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 41.4;
Pred. No. 2e
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
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                                                                                                                                                                                                                                                                                            gene therapy;
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밁 8

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AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide sequences (I), which contain single nucleotide polymorphisms (SNPs).

CC AAM53114 to AAM5329 represent peptides related to human polymorphic contain single nucleotide polymorphisms (SNPs).

CC polynucleotide sequences. The sequences can be used in gene and protein contempy, and in vaccine production. (I) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases cassociated with inappropriate expression of polymorphic polypeptides. For caxemple, (I) may be used to treat disorders by rectifying mutations or caxemple, (I) may be used to treat disorders by rectifying mutations or caxemple, (I) may be used to treat disorders by rectifying mutations or compression of polypeptides. Additionally, (I) and its complementary confidences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and confidence therefore which patients may be in need of restorative therapy. The contibodies specific for polymorphic polypeptides. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic colleges may also be used as diagnostic agents for detecting the presence of polymorphic colleges may also be used as diagnostic agents for detecting the presence of polymorphic colleges may also be used as diagnostic agents for detecting the presence of polymorphic colleges may also be used as diagnostic agents for detecting the presence of polymorphic colleges may also be used as diagnostic agents for detecting the presence of polymorphic colleges may also be used as diagnostic agents for detecting the presence of polymorphic colleges may also be used as diagnostic agents for detecting the presence of polymorphic colleges may also be used as diagnostic agents for detecting the presence of polymorphic colleges may also be used as diagnostic agents for detecting the presence of polymorphic colleges may also be used as diagn
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Sequence 51 BP; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Polymorphic nucleic acid sequences, useful in genetic testing and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Page
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    A; 14 C; 11 G; 9
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        T; 0 U; 0 Other;
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Query Match
Best Local S
Matches 45
              1085 TAGAGGCGGGGTTTCACCATATTTGTCAGGCTGGTCTCAAACTCCTGACCT 1135
51 TAGAGACGGGGTTTCACCATTTTGGTTAGGCTGGTCTTGAACTCCTGACCT 1
                                              45;
                                                         Similarity
                                              Conservative
                                                         4.2%;
                                              <u>,</u>
                                                           Score 41.4;
Pred. No. 2
                                                 Mismatches
                                                           4; DB 1;
2e+02;
                                                 6
                                                                       Length
                                                  Indels
                                                 0,
                                                  Gaps
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RESULT 131
AAI79539/c
ID AAI795
AAI79539 standard; DNA;
                                                   51
                                                   ВP
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Human silent SNP containing nucleic acid SEQ:6480 09-NOV-2001 (first entry)

Human; single nucleotide polymorphism; SNP; genome; gene therapy; protein therapy; vaccine; probe; diagnostic assay; detection; quantitation; restorative therapy; polymorphic; ds.

Homo sapiens.

WO200140521-A2

07-JUN-2001.

30-NOV-2000; 2000WO-US032758

30-NOV-1999; 99US-0168138P 29-NOV-2000; 2000US-00726173

(CURA-) CURAGEN CORP.

Shimkets RA, Leach M;

WPI; 2001-356160/37

Polymorphic nucleic acid sequences, useful ij genetic testing and

Claim 1; Page 2490; 2653pp; English

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CC sequences (I), which contain single nucleotide polymorphic polynucleotide CR sequences (I), which contain single nucleotide polymorphisms (SNPs). CC AAM53114 to AAM53329 represent peptides related to human polymorphic CR cambility, and in vaccine production. (I) and the polypeptides encoded by therapy, and in vaccine production. (I) and the polypeptides encoded by the may be used in the prevention, diagnosis and treatment of diseases CC associated with inappropriate expression of polymorphic polypeptides. For CC deletions in a patient's genome that affect the activity of polypeptides or CC deletions in a patient's genome that affect the patients own CC production of polypeptide. Additionally, (I) and its complementary CC sequences may also be used as DNA probes in diagnostic assays to detect CC and quantitate the presence of similar nucleic acids in samples, and CC therefore which patients may be in need of restorative therapy. The CC continuous production of polypeptides encoded by (I) may be used as antigens in the production of CC polypeptides encoded by (I) may be used as antibodies may also be used to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic
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Best Local S
Matches 45
AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide sequences (1), which contain single nucleotide polymorphisms (SNPs).

AAM53114 to AAM53329 represent peptides related to human polymorphic polynucleotide sequences. The sequences can be used in gene and protein therapy, and in vaccine production. (1) and the polypeptides encoded by
                                                                                                                                                                                        Claim 1;
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29-NOV-2000; 2000US-00726173
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                                                                                                                                                                                                                                                                                 Polymorphic nucleic acid sequences, useful in genetic testing
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Pred. No. 2e+02;
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RESULT 133
AAI76092/c
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AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide sequences (I), which contain single nucleotide polymorphisms (SNPs). AAM53114 to AAM53129 represent peptides related to human polymorphic polymucleotide sequences. The sequences can be used in gene and protein therapy, and in vaccine production. (I) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of polymorphic polypeptides. For deletions in a patient's genome that affect the activity of polypeptides by expressing inactive proteins or to supplement the patients own production of polypeptide. Additionally, (I) and its complementary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    associated with inappropriate expression of polymorphic polypeptides. For example, (I) may be used to treat disorders by rectifying mutations or deletions in a patient's genome that affect the activity of polypeptides by expressing inactive proteins or to supplement the patients own production of polypeptide. Additionally, (I) and its complementary sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and therefore which patients may be in need of restorative therapy. The polypeptides encoded by (I) may be used as antigens in the production of antibodies specific for polymorphic polypeptides. The antibodies may also be used to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic polyportides.
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29-NOV-2000; 2000US-00726173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; single nucleotide polymorphism; SNP; genome; protein therapy; vaccine; probe; diagnostic assay; quantitation; restorative therapy; polymorphic; ds.
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CC AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide CC sequences (I), which contain single nucleotide polymorphisms (SNPs). CC AAM53114 to AAA5329 represent peptides related to human polymorphic CC polynucleotide sequences. The sequences can be used in gene and protein CC therapy, and in vaccine production. (I) and the polypeptides encoded by CC them may be used in the prevention, diagnosis and treatment of diseases CC associated with inappropriate expression of polymorphic polypeptides. For CC example, (I) may be used to treat disorders by rectifying mutations or CC example, (I) may be used to treat disorders by rectifying mutations or CC eletions in a patient's genome that affect the activity of polypeptides by expressing inactive proteins or to supplement the patients own CC production of polypeptide. Additionally, (I) and its complementary CC sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and CC therefore which patients may be in need of restorative therapy. The CC antibodies specific for polymorphic polypeptides. The antibodies may also be used as antigens in the production of antibodies specific for polymorphic polypeptides. The antibodies may also
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Best Local S
Matches 45
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAI79838 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   polypeptides in samples
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30-NOV-1999; 99US-0168138P.
29-NOV-2000; 2000US-00726173.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 1; Page 2579; 2653pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Polymorphic nucleic acid sequences, useful in genetic testing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-NOV-2000; 2000WO-US032758.
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restorative therapy; polymorphic; ds.
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                                                   continuous control of the polymenton of the polymenton of the polymenton of diseases content may be used in the prevention, diagnosis and treatment of diseases cassociated with inappropriate expression of polymorphic polymeptides. For cexample, (I) may be used to treat disorders by rectifying mutations or cexample, (I) may be used to treat disorders by rectifying mutations or complete the activity of polymeptides on the prevention of polymeptides on the production of polymeptides or to supplement the patients own composition of polymeptides or production of polymeptides. Additionally, (I) and its complementary completed the presence of similar nucleic acids in samples, and confident the production of contibodies may also be used as antigens in the production of colymeptides encoded by (I) may be used as antigens in the production of contibodies specific for polymorphic polymeptides. The antibodies may also be used to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic colymeptides in samples
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide sequences (I), which contain single nucleotide polymorphisms (SNPs). AAM53114 to AAM53329 represent peptides related to human polymorphic
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C AAM53114 to AAM5329 represent peptides related to human polymorphic polypeptides. For certain may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of polymorphic polypeptides. For certain polymorphic polypeptides. For certain complex polypeptides for certain polymorphic polypeptides for certain polymorphic polypeptides. For certain polymorphic polypeptides or complement the patients own certain polymorphic polypeptides or complement the patients own certain polymorphic poly
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; single nucleotide polymorphism; SNP; genome; gene therapy; protein therapy; vaccine; probe; diagnostic assay; detection; quantitation; restorative therapy; polymorphic; ds.
                                                                                                                                                                         Sequence 51 BP; 10 A; 12 C; 18 G; 11 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Polymorphic nucleic acid sequences, useful in genetic testing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30-NOV-1999; 99US-0168138P.
29-NOV-2000; 2000US-00726173.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAI79697 standard; DNA; 51 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (CURA-)
                                       1025 CCCAAGCAGCTGGGATTACGGGCACCTGCCACCACACCCCGCTAATTTTTG 1075
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      692 TCCCGGGTTCAAGTTATTCTCCTGCCCCAGCCTCCTGAGTAGCTGGGACTA 742
51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           51
                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ₽Ą,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Page 2537; 2653pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ilarity 88.2%;
Conservative
                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Leach M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     amino acid changing SNP nucleic acid SEQ:6638
                                                                                                       4.2%;
                                                                                  Score 41.4; DB 1
Pred. No. 2e+02;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0,
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0; Mismatches
                                                                                                                              1;
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                                                                                                                              Length 51;
                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
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                                                                                    Gaps
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RESULT 138 AAI73250/c ID AAI732 XX

AAI73250 standard; DNA;

51 BP

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847 CCTCGGCCTCCCAAAGTGCTGGGATTACAGGCGTGAGCCACCACGCCCGGC 897 CCTCGGCCTCCCAAATTCCTGGGACTACAGGCGTGAGCCACTGCACCCGGC

<u>,,</u>

Mismatches

6

Indels

0;

0

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RESULT 137
AAI74778
ID AAI7477
XX AAI747
XX AAI747
XX Human;
XX Human;
XX Human;
XX Homo s
XX Homo s
XX Homo s
XX WO2001
XX 30-NOV
XX 30-NOV
XX 30-NOV
XX 30-NOV
XX 30-NOV
XX AOINS
PR WPI; 2
XX PPlymc
PR WPI; 2
XX AAI731
CC Sequer
CC AAM531
CC Sequer
CC ASSocio
CC Sequer
                                                                                                                                                                     CC AANT3166 to AANT9867 represent isolated human polymorphic polynucleotide CC sequences (I), which contain single nucleotide polymorphisms (SNPs). CC AAM53114 to AAM53329 represent peptides related to human polymorphic CC polynucleotide sequences. The sequences can be used in gene and protein CC therapy, and in vaccine production. (I) and the polypeptides encoded by C them may be used in the prevention, diagnosis and treatment of diseases CC associated with inappropriate expression of polymorphic polypeptides. For CC example, (I) may be used to treat disorders by rectifying mutations or CC example, (I) may be used to treat ffect the activity of polypeptides or CC by expressing inactive proteins or to supplement the patients own CC production of polypeptide. Additionally, (I) and its complementary CC sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and CC polypeptides encoded by (I) may be used as antigens in the production of contibodies specific for polymorphic polypeptides. The antibodies may also be used to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic colypeptides in samples in samples.
Query Match
Best Local Similarity
Matches 45; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    30-NOV-1999; 99US-0168138P.
29-NOV-2000; 2000US-00726173.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    09-NOV-2001
                                                                                                                        Sequence 51
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-356160/37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Shimkets RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-NOV-2000; 2000WO-US032758
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Polymorphic nucleic acid sequences, useful in genetic testing and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (CURA-) CURAGEN CORP
                                                                                                                        BP; 9 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Leach M;
                                 4.2%;
                                                                                                                        21 C; 13 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      51
                                    Score 41.4; DB 1; Length 51; Pred. No. 2e+02;
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AAI79700/c
ID AAI797
XX
AC AAI797
XX
DT 09-NOV
XX
DE Human
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CC AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide contents of the polynorphic polynucleotide sequences (I), which contain single nucleotide polymorphisms (SNPs). CC AAM53114 to AAM53329 represent peptides related to human polymorphic polynucleotide sequences. The sequences can be used in gene and protein therapy, and in vaccine production. (I) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases composed the production of polymorphic polypeptides. For example, (I) may be used to treat disorders by rectifying mutations or completions in a patient's genome that affect the activity of polypeptides for production of polypeptide. Additionally, (I) and its complementary composed by expressing inactive proteins or to supplement the patients own composed by the patients of the sequences may also be used as DNA probbes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and complements the production of polypeptides may be in need of restorative therapy. The complement of complements of the polypeptides of the production of complements may be used as antigens in the production of complements 
                                                                                                                                                                                                               RESULT 139
                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
Matches 45; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; single nucleotide polymorphism; SNP; genome; gene therapy; protein therapy; vaccine; probe; diagnostic assay; detection; quantitation; restorative therapy; polymorphic; ds.
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29-NOV-2000; 2000US-00726173
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Polymorphic nucleic acid sequences, useful in genetic testing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2001-356160/37
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 51 BP; 11 A; 13 C; 18 G; 9 T; 0 U; 0 Other;
                           Human conservative amino acid changing
                                                                       09-NOV-2001
                                                                                                                                                                   AAI79700 standard; DNA; 51 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           polypeptides in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                 956 GCAATGGCCAAATCTCGGCTCACTGCAACCTCTGCCTCCCGGGCTCAAGCG 1006
                                                                                                                                                                                                                                                                                     51 GCAGTGGCATGATCTTGGCTCACTGCAACCTCTGCCTCCCAGGCTCAAGCG 1
                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SNP containing nucleic acid SEQ:191.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
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                                                                          (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           samples
                                                                                                                                                                                                                                                                                                                                                                                                       4.2%;
                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                       Score 41.4; DB
Pred. No. 2e+02;
                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                             SNP nucleic acid SEQ:6641.
                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1;
                                                                                                                                                                                                                                                                                                                                                                                   6,
                                                                                                                                                                                                                                                                                                                                                                                                                               Length 51;
                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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SXFXFXBXFXR

Homo sapiens

Human; single nucleotide polymorphism; SNP; genome; gene therapy; protein therapy; vaccine; probe; diagnostic assay; detection; quantitation; restorative therapy; polymorphic; ds.

Human silent SNP containing nucleic acid SEQ:5327.

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CC polynciectide sequences. The sequences can be used in gene and protein content may be used in the prevention, (i) and the polypeptides encoded by caseociated with inappropriate expression of polymorphic polypeptides. For content can be used to treat disorders by rectifying mutations or content in a patient's genome that affect the activity of polypeptides. For conduction of polypeptide. Additionally, (i) and its complementary confidences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and therefore which patients may be used as antigens in the production of polypeptides. For confidence specific for polymorphic polypeptides. The antibodies may also be used as diagnostic agents for detecting the presence of similar nucleic acids in samples, and contibodies specific for polymorphic polypeptides. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic polypeptides. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic polypeptides may also be used as diagnostic agents for detecting the presence of polymorphic polypeptides in samples may also be used as diagnostic agents for detecting the presence of polymorphic polypeptides in samples.
                                                                                                                                                                                                     RESULT 140
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   quantitation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sequences (I), which contain single nucleotide polymorphisms waves, AAMS3114 to AAMS3329 represent peptides related to human polymorphic AAMS3114 to AAMS3329 represent peptides related to human polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAI73060 to AAI79867 represent isolated human polymorphic pol
sequences (I), which contain single nucleotide polymorphisms
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1; Page 2538; 2653pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2001-356160/37.
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29-NOV-2000; 2000US-00726173.
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   09-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 51 BP; 10 A; 12 C; 17 G; 12 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Polymorphic nucleic acid sequences, useful in genetic testing and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (CURA-) CURAGEN CORP.
                                                                                                                                       AAI78386 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local
                                                                                                                                                                                                                                                                                                                                                           836 TGATCTGCCTGCCTCGGCCTCCCAAAGTGCTGGGATTACAGGCGTGAGCCA 886
                                                                                                                                                                                                                                                                                                                  51 TGATCCGCCCATCTCGGCCTCCCAAAATGCTGGGATTACAGGCATGAGCCA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                         45;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Leach M;
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 41.4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             6,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 51;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
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CC AAM53114 to AAM5329 represent peptides related to human polymorphic polymo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Loca
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                              Human; single nucleotide polymorphism; SNP; genome; gene therapy; protein therapy; vaccine; probe; diagnostic assay; detection; quantitation; restorative therapy; polymorphic; ds.
                    30-NOV-2000; 2000WO-US032758.
                                                                    07-JUN-2001.
                                                                                                                                                                                                                                                                                                               Human silent SNP containing nucleic acid SEQ:803.
                                                                                                                                                                                                                                                                                                                                                                    09-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAI73862 standard; DNA; 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 51 BP; 9 A; 17 C; 16 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AA173060 to AA179867 represent isolated human polymorphic polynucleotide sequences (I), which contain single nucleotide polymorphisms (SNPs).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; Page 2140; 2653pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Polymorphic nucleic acid sequences, useful in genetic testing and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2001-356160/37
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      30-NOV-1999;
29-NOV-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        646
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45; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AGGCTGGAGTGCAGTGGCGCAATCTTGGCTCACTGCAACCTCTGCCTCCCG 696
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2000US-00726173.
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                                                                                                                                                                                                                                                                                                                                                                  (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 41.4; DB 1; Length 51; Pred. No. 2e+02; o; Mismatches 6; Indels
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RESULT 142
AA179783/c
ID AA1797
XX AA1797
XX AA1797
XX O9-NOV
XX Human
XX Human;
XW Quanti
XX Quanti
XX WO2001
XX WO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; single nucleotide polymorphism; SNP; genome; protein therapy; vaccine; probe; diagnostic assay; quantitation; restorative therapy; polymorphic; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             polypeptides in
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30-NOV-1999; 99US-0168138P.
29-NOV-2000; 2000US-00726173.
                                                                                                                        30-NOV-1999; 99US-0168138P.
29-NOV-2000; 2000US-00726173.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human nonconservative amino acid changing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     09-NOV-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 51 BP; 10 A; 21 C; 8 G; 12 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1; Page 300; 2653pp; English
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                                                                                                                                                                                                                            30-NOV-2000; 2000WO-US032758
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         671 TGGCTCACTGCAACCTCTGCCTCCCGGGTTCAAGTTATTCTCCTGCCCCAG 721
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         45;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    samples
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 41.4; I
Pred. No. 2e+0
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2e+02; Length 51;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SNP nucleic acid SEQ:6724.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; gene therapy;
detection;
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Shimkets RA,

(CURA-) CURAGEN CORP.

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RESULT 143
AAH90585/c
ID AAH905
XX AAH905
XX AAH905
XX Human
XX Human;
KW Gene t
XX Homo :
XX Ho
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AAM53314 to AAM53329 represent peptides related to human polymorphic polymorphic polymorphic polymorphic polymorphic polymorphic polymorphic states and the sequences. The sequences can be used in gene and protein therapy, and in vaccine production. (I) and the polymorphic polymorphic consociated with inappropriate expression of polymorphic po
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Best Local S
Matches 45
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human clone cg43080072 SNP site, SEQ ID NO:465
                                      WPI; 2001-425617/45.
                                                                                                                                                                                                                                                                                                                                                                                                             05-JUL-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200147942-A2
                                                                                                                Shimkets RA, Leach M;
                                                                                                                                                                                                                                                                 27-DEC-1999;
                                                                                                                                                                                                                                                                                                                                       27-DEC-2000; 2000WO-US035387
                                                                                                                                                                                          (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  989 GCCTCCGGGCTCAAGCGATTCTCCTGTCTCAGCCTCCCAAGCAGCTGGGA 1039
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11 Similarity
45; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 nucleotide polymorphism; SNP; detection; identification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  genetic
                                                                                                                                                                                                                                                                 99US-00472865
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers replace(26,C)
/*tag= a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             'standard_name= "single nucleotide polymorphism"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              disorder;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
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cc pairs of nucleotides containing SNPs which result in changes in the corresponding amino acid sequences (AAG64751-AAG64762). The SNPs in cc corresponding amino acid sequences (AAG64751-AAG64762). The SNPs in cc sequences 574 (AAH90699-AAH90694) lead to conservative amino acid cc changes, while those in sequences 575 to 578 (AAH90695-AAH90698) result cr non-conservative changes. The SNP in sequences 579 and 580 (AAH90699-AAH90700) generates a frameshift mutation. The invention also relates to call a method of detercting a polymorphic site in a nucleic acid and a method cc of determining the relatedness of two nucleic acids. It also encompasses peptides containing polymorphic sites, antibodies raised against such competities, and a method of detecting polymorphic proteins/ peptides using the antibodies. The nucleic acids are useful for gene therapy of an conditional having, suspected of having, or at risk of developing a condition due to the presence of a sequence polymorphism. Such treatment would comprise administration of the wild-type nucleic acid sequence. Antibodies raised against polymorphic peptides can also be considered as a sequence polymorphism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequences AAH90121-AAH90168 polymorphisms (SNPs). Sequences 1 to 568 contain single nucleotide polymorphisms of nucleotides which contain (AAH90121-AAH90688) are consecutive pairs of nucleotides which contain silent SNPs. Sequences 569 to 580 (AAH90689-AAH90700) are consecutive silent SNPs. Sequences 569 to 580 (AAH90689-AAH90700) are consecutive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequences AAH90121-AAH90700 represent 580 human cDNA sequences contain single nucleotide polymorphisms (SNPs). Sequences 1 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1; Page 109;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New polynucleotides containing single nucleotide polymorphisms, detecting the presence of polymorphism, detecting a polymorphic treating a patient suffering from a pathology ascribed to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  295pp; English.
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RESULT 144
                                                                    Query Match
Best Local S
Matches 45
                                                                                                                              Sequence 51 BP; 12 A; 14 C; 15 G; 10 T; 0 U; 0 Other;
                                                       836 TGATCTGCCTGGCCTCGGCCTCCCAAAGTGCTGGGATTACAGGCGTGAGCCA 886
                                   51 TGATCCACTCGCTTCGGCCTCCCAAAGTGCTGGGATTATAGGCGTGAGCCA 1
                                                                                45;
                                                                                             Similarity
                                                                                Conservative
                                                                                            4.2%;
                                                                                0
                                                                                             Score 41.4;
Pred. No. 26
                                                                                   Mismatches
                                                                                              2e+02;
                                                                                                         DB
                                                                                                         1;
                                                                                   6;
                                                                                                        Length
                                                                                    Indels
                                                                                   0
                                                                                    Gaps
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in the treatment of such individuals

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Human; single nucleotide polymorphism; SNP; forensic test; aberrant protein expression;
                                                                                                                                            Human coding sequence polymorphic site SEQ ID NO:
                                                                                                                                                         01-OCT-2001
                                                                                                                                                                     AAH89405;
                                                                                                                                                                                AAH89405 standard; DNA;
                                                                                                     WO200151670-A2
                                                                                                                Homo sapiens
                                                                                                                                                          (first entry)
                                                                                                                                                                                51
                                                                                                                                                                                ВP
                                                                                                                             paternity test; ds.
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05-JAN-2001; 2001WO-US000322

19-JUL-2001.

07-JAN-2000; 2000US-0174962P

CURAGEN CORP.

Isolated human polynucleotides containing single nucleotide polymorphisms, useful for the treatment and diagnosis of e.g. cancer.

diabetes

WPI; 2001-451871/48. P-PSDB; AAM00292.

Shimkets RA,

Leach

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8 × 55 55 55 55 55 55 55 55 55 8
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AAH89485/c
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Best Local S
Matches 45
          The present invention relates to human nucleic acids containing single nucleotide polymorphisms (SNPs). These can be used in forensic and paternity tests, and to aid in the treatment of diseases associated with aberrant protein expression, including cancer, amyloidosis, diabetes, Alzheimer's disease, Down's syndrome, oedema, lupus (SLE), vasculitis, glomerulonephritis, haemolytic anaemia, thrombocytopaenia, arthritis, meningitis, muscular disorders, dementia, neurological diseases, tuberous sclerosis, male infertility, hypercalcaemia, blood pressure disorders, osteoporosis, pathogenic infections, hypercholesterolaemia, obesity or autoimmunity. The present sequence is a polymorphism-containing oligonucleotide fragment of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   nucleotide polymorphisms (SNPs). These can be used in forensic and paternity tests, and to aid in the treatment of diseases associated with aberrant protein expression, including cancer, amyloidosis, diabetes, Alzheimer's disease, Down's syndrome, oedema, lupus (SLE), vasculitis, glomerulonephritis, haemolytic anaemia, thrombocytopaenia, arthritis, menningitis, muscular disorders, dementia, neurological diseases, tuberous sclerosis, male inferrility, hypercalcaemia, blood pressure disorders, osteoporosis, pathogenic infections, hypercholesteroleemia, obesity or autoimmunity. The present sequence is a polymorphism-containing
                                                                                                                                                                                                                                                           Isolated human polynucleotides contain: polymorphisms, useful for the treatment infection and diabetes.
                                                                                                                                                                                                                                                                                                                                   WPI; 2001-451871/48.
P-PSDB; AAM00370.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; single nucleotide polymorphism; SNP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human coding sequence polymorphic site
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    forensic test;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 present invention relates to human nucleic acids containing leotide polymorphisms (SNPs). These can be used in forensic a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   663 CGCAATCTTGGCTCACTGCAACCTCTGCCTCCCGGGTTCAAGTTATTCTCC 713
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                                                                                                                                                                                                                                                                                                                                                                                                                       CURAGEN CORP
                                                                                                                                                                                                                                                                                                                                                                                    RA,
                                                                                                                                                                                                                            Page 180; 475pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CACGATCTTGGCTCACTGCAACCTCCGCCTCCCAGGTTCAAGTGATCCTCC 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Page 159; 475pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                    Leach MD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   aberrant protein expression;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 41.4; DB : Pred. No. 2e+02;
                                                                                                                                                                                                                                                                             containing single nucleotide treatment and diagnosis of e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    paternity test; ds.
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RESULT 146
AAH89514/c
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Best Local Similarity
Matches 45; Conserv
                                                      Matches
                                                                   Query Match
Best Local Similarity
                                                                                                                                   The present invention relates to human nucleic acids containing single nucleotide polymorphisms (SNPs). These can be used in forensic and paternity tests, and to aid in the treatment of diseases associated with aberrant protein expression, including cancer, amyloidosis, diabetes, Alzheimer's disease, Down's syndrome, oedema, lupus (SLE), vasculitis, glomerulonephritis, haemolytic anaemia, thrombocytiopaenia, arthritis, meningitis, muscular disorders, dementia, neurological diseases, tuberous sclerosis, male infertility, hypercalcaemia, blood pressure disorders, osteoporosis, pathogenic infections, hypercholesterolaemia, obesity or autoimmunity. The present sequence is a polymorphism-containing oligonucleotide fragment of the invention
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                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                              Claim 1;
                                                                                                                                                                                                                                                                                                                                                       Isolated human polynucleotides containing single nucleotide polymorphisms, useful for the treatment and diagnosis of e.g. cancer,
                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2001-451871/48.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                07-JAN-2000; 2000US-0174962P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; single nucleotide polymorphism; SNP; forensic test; aberrant protein expression;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human coding sequence polymorphic site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAH89514 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                  P-PSDB; AAM00397.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200151670-A2
                                                                                                                                                                                                                                                                                                                                       infection and diabetes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (CURA-) CURAGEN CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1080 TTCATTAGAGGCGGGTTTCACCATATTTGTCAGGCTGGTCTCAAACTCCT 1130
                          844 CTGCCTCGGCCTCCCAAAGTGCTGGGATTACAGGCGTGAGCCACCACGCCC 894
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            51
51
                                                      45;
                                                                                                             51
                                                                                                                                                                                                                                                                                                            Page 188; 475pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TTTAGTAGAGACAGGGTTTCACCATATTGGCCAGGCTGGTCTCAAACTCCT
 BP; 7 A;
                                                     Conservative
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                                                                                                             13 C;
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                                                                   4.2%;
88.2%;
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                                                                                                             22 G; 9 T; 0 U; 0 Other;
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                                                                   Score 41.4; DB 1;
Pred. No. 2e+02;
                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           paternity test; ds.
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                                                                                 Length
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                                                        Indels
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RESULT 147 AAH89519/c ID AAH89519 :

standard; DNA;

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RESULT 148
AAH99467/c
ID AAH894
XX AAH894
XX AAH894
XX Human
XX Human
XX Human
XX Human
XX Horens
XX Homo 8
XX Homo 8
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Best Local Similarity
Matches 45; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        07-JAN-2000; 2000US-0174962P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; single nucleotide polymorphism; SNP; forensic test; aberrant protein expression;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO200151670-A2
                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 51 BP; 10 A; 16 C; 16 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Isolated human polynucleotides containing single nucleotide polymorphisms, useful for the treatment and diagnosis of e.g. cancer, infection and diabetes.
                                                                      Human; single nucleotide polymorphism; SNP; forensic test; aberrant protein expression;
                                                                                                               Human coding sequence
                                                                                                                                             01-OCT-2001
                                                                                                                                                                           AAH89467;
                                                                                                                                                                                                       AAH89467 standard; DNA; 51 BP
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                WO200151670-A2
                                           Homo sapiens.
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                                                                                                                                                                                                                                                                                                            GTGATCTGCCTGGCCTCGGCCTCCCAAAGTGCTGGGATTACAGGCGTGAGCC 885
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                                                                                                                polymorphic site
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                                                                                                                                                                                                                                                                                                                                                         Score 41.4;
Pred. No. 26
                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                         2e+02;
                                                                                                                  SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                       DB 1;
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                                                                          paternity test; ds.
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AAH89553/c
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  RESULT 149
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Best Local S
Matches 45
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to human nucleic acids containing single nucleotide polymorphisms (SNPs). These can be used in forensic and paternity tests, and to aid in the treatment of diseases associated with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 1; Page 176; 475pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               07-JAN-2000; 2000US-0174962P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 51 BP; 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Isolated human polynucleotides containing single nucleotide polymorphisms, useful for the treatment and diagnosis of e.g. cancer, infection and diabetes.
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                                                                                                                                                                                                                                                        Human; single nucleotide polymorphism; SNP; forensic test; aberrant protein expression;
                                                                                                                                                                                                                                                                                                 Human
                                                                                                                                                                                                                                                                                                                           01-OCT-2001
                                                                                                                                                                                                                                                                                                                                                      AAH89553;
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                                                                Shimkets RA,
                                                                                                                     07-JAN-2000; 2000US-0174962P
                                                                                                                                             05-JAN-2001; 2001WO-US000322
                                                                                                                                                                                                     WO200151670-A2
                                                                                                                                                                                                                              Homo sapiens
Isolated human polynucleotides containing single nucleotide
                                        WPI; 2001-451871/48
                                                                                                                                                                          19-JUL-2001.
                                                                                             (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                               coding
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TCTTGATCTCTGGACCTTGTGATCTGCCTGCCTCGGCCTCCCAAAGTGCTG 867
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                              sequence polymorphic site SEQ ID NO: 334.
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                                                                  Leach MD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; 10 C; 21 G; 7 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 41.4; DB 1
Pred. No. 2e+02;
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                                                                                                                                                                                                                                                           paternity test; ds.
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Best Local
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The present invention relates to human nucleic acids containing single nucleotide polymorphisms (SNPs). These can be used in forensic and paternity tests, and to aid in the treatment of diseases associated with aberrant protein expression, including cancer, amyloidosis, diabettes, Alzheimer's disease, Down's syndrome, oedema, lupus (SLE), vasculitis, glomerulonephritis, haemolytic anaemia, thrombocytopaenia, arthritis, meningitis, muscular disorders, dementia, neurological diseases, tuberous sclerosis, male infertility, hypercalcaemia, blood pressure disorders, osteoporosis, pathogenic infections, hypercholesterolaemia, obesity or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               aberrant protein expression, including cancer, amyloidosis, diabetes, Alzheimer's disease, Down's syndrome, oedema, lupus (SLE), vasculitis, glomerulonephritis, haemolytic anaemia, thrombocytopaenia, arthritis, meningitis, muscular disorders, dementia, neurological diseases, tuberous sclerosis, male infertility, hypercalcaemia, blood pressure disorders, osteoporosis, pathogenic infections, hypercholesterolæmia, obesity or autoimmunity. The present sequence is a polymorphism-containing oligonucleotide fragment of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to human nucleic acids containing single nucleotide polymorphisms (SNPs). These can be used in forensic and paternity tests, and to aid in the treatment of diseases associated with aberrant protein expression, including cancer amplifications of the containing single paternal protein expression, including cancer amplifications.
                                                                                                                                                                   Claim 1;
                                                                                                                                                                                                                Isolated human polynucleotides containing single nucleotide polymorphisms, useful for the treatment and diagnosis of e.g. cancer,
                                                                                                                                                                                                                                                                                                                                                                                    07-JAN-2000; 2000US-0174962P
                                                                                                                                                                                                                                                                                                                                                                                                                   05-JAN-2001; 2001WO-US000322.
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DB; AAM00357.
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                                                                                                                                                                                                                                                                                                                                                     CURAGEN CORP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP; 14 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
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                                                                                                                                                                                                      diabetes
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Pred. No. 2e+02;
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expression;
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                                                                                                                                                              This invention relates to a novel isolated protein which comprises a human mannosyl transferase having the same sequence as the fully define 611- or 255-amino acid sequence or its fragment. The invention may be useful for the production of compounds with an antimanic or antidepressant activity whilst the disclosed sequences may be used for gene therapy. The invention also provides a human mannosyl transferase fusion protein and a chromosome 9 fusion protein, both of which result from a chromosome 11 translocation. The human mannosyl transferase and the fusion proteins are useful for diagnosing or predicting the susceptibility to a bipolar disorder and for identifying a compound that of a region of human DNA surrounding a single nucleotide polymorphism within the gene which encodes the human mannosyl transferase.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New polypeptide comprising human mannosyl transferase, useful for diagnosing or predicting the susceptibility to a bipolar disorder identifying a compound that modulates the activity of a mannosyl
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Pred. No. 26
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                                                                                                                                                           AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide consequences (I), which contain single nucleotide polymorphisms (SNPs). CC AAM53114 to AAM5329 represent peptides related to human polymorphic polynucleotide sequences. The sequences can be used in gene and protein cc therapy, and in vaccine production. (I) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases cc associated with inappropriate expression of polymorphic polypeptides. For cc example, (I) may be used to treat disorders by rectifying mutations or cc example, (I) may be used to treat disorders by rectifying mutations or cc productions in a patient's genome that affect the activity of polypeptides by expressing inactive proteins or to supplement the patients own cc production of polypeptide. Additionally, (I) and its complementary csequences may also be used as May probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and ctherefore which patients may be in need of restorative therapy. The polypeptides encoded by (I) may be used as antigens in the production of complementary complements of the used to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic colleges in camples.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human silent SNP containing nucleic acid SEQ:6530.
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29-NOV-2000; 2000US-00726173
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; Page 2505; 2653pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Polymorphic nucleic acid sequences, useful in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-356160/37
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CCTGCCTCGGCCTCCCAAAGTGCTGGGATTACAGGCGTGAGCCACC 888
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Pred. No. 2.1e-
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RESULT 154
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र
                                                                                                                                                                                                                                                                                                                          present invention describes newly discovered deletion mutations that are believed to be deleterious and cause significant alterations in the structure or biochemical function of BRCA1. Accordingly, it provides methods for detecting such mutants, as well as identifying and screening for cytostatic compounds useful for treating or preventing cancers associated with a BRCA1 genetic variant. This polymucleotide is a mutant human BRCA1 genomic DNA fragment that arises as a result of a recombination event (deletion 1), which causes the omission of exons 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This invention relates to a novel method for predicting a predisposition to cancer in a patient by detecting large deletions in the human tumour suppressor gene identified as the breast cancer susceptibility gene 1 (BRCA1). Specifically, it refers to deletions that result from the unequal crossover between a pair of repetitive Alu sequences in the BRCA1 gene, such that the recombined nucleotide sequence containing the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Predicting a predisposition to cancer in a patient comprising detecting deletion in the BRCA1 gene that results from the unequal crossover between a pair of repetitive sequences in the BRCA1 gene.
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09-AUG-2002; 2002US-0402430P
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breast cancer susceptibility gene 1; BRCAl; repetitive Alu;
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                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                    recombination event
and 17, given in an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               deletion indicates a predisposition to breast and ovarian cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; SEQ ID NO 15; 59pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2004-062369/06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Scholl T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             09-JUN-2003; 2003WO-US018098
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (MYRI-) MYRIAD GENETICS INC
                                                            843
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4 CCCACCTCGGCCTCCCAAAGTGCTGGGATTACAGGTGTGAGCCACC
                                                                                                                        44;
1 CCCGTCTCGGCCTCCCAAAGTGCTGGGATTACAGGTGTGAGCCATCGCG
                                                                                                                                                                                                                                                                                                    given in an exemplification of the
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                                      CTGCCTCGGCCTCCCAAAGTGCTGGGATTACAGGCGTGAGCCACCACG
                                                                                                                                                                                                                                              49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hendrickson
                                                                                                                                                                                                                                           BP;
                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                       8 A; 16 C; 15 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        genomic DNA resulting
                                                                                                                                                     4.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BC, Ward B,
                                                                                                                           0
                                                                                                                                                        Score 41; DB 1;
Pred. No. 2.1e+02;
                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pruss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           from
                                                                                                                                                                                                                                                                                                             invention.
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                                                                                                                           5
                                                                                                                                                                                 Length 49;
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                                                                                                                               Gaps
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RESULT 155
AAI79093/C
ID AAI790
XX
AC AAI790
XX
DT 09-NOV
DT 09-NOV
XX
XX
XX
XX
KW Human;
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                                                                                                                                                                                                                                                                                                                                                                                                                      AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide sequences (I), which contain single nucleotide polymorphisms (SNP8). CC AAM53124 represent peptides related to human polymorphic polymorphic cc polynucleotide sequences. The sequences can be used in gene and protein cc therapy, and in vaccine production. (I) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases cc associated with inappropriate expression of polymorphic polypeptides. For cc example, (I) may be used to treat disorders by rectifying mutations or cc example, (I) may be used to treat affect the activity of polypeptides by expressing inactive proteins or to supplement the patients own cc production of polypeptide. Additionally, (I) and its complementary cc sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and ct therefore which patients may be in need of restorative therapy. The polypeptides encoded by (I) may be used as antigens in the production of antibodies specific for polymorphic polypeptides. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic polypeptides in samples and activity. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic polypeptides in samples.
                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                    Matches
 Human; single nucleotide polymorphism; SNP; genome; gene therapy;
                                    Human silent SNP containing nucleic acid SEQ:6034.
                                                                            09-NOV-2001
                                                                                                                                                 AAI79093 standard;
                                                                                                                                                                                                                                                                                                                                                                                           Sequence 51 BP; 8 A; 16 C; 13 G; 14 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; Page 1200; 2653pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30-NOV-1999; 99US-0168138P.
29-NOV-2000; 2000US-00726173.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30-NOV-2000; 2000WO-US032758
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; single nucleotide polymorphism; SNP; genome; gene therapy; protein therapy; vaccine; probe; diagnostic assay; detection; quantitation; restorative therapy; polymorphic; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           09-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Polymorphic nucleic acid sequences, useful in genetic testing and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                      Local
                                                                                                                                                                                                                                                                                193 TTCTCCATGTTGGTCAGGCTGGTCTCGAACTCCCGACCTCAGATGATCC
                                                                                                                                                                                                                                                                                                                  44;
                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                    Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Leach M;
                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             99US-0168138P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      containing nucleic acid SEQ:3757.
                                                                                                                                                 DNA;
                                                                                                                                                                                                                                                                                                                                    4.1%;
                                                                                                                                                   51
                                                                                                                                                                                                                                                                                                                    0; · Mismatches
                                                                                                                                                                                                                                                                                                                                      Score 41;
Pred. No.
                                                                                                                                                                                                                                                                                                                                       2.1e+02;
                                                                                                                                                                                                                                                                                                                                                         DB 1;
                                                                                                                                                                                                                                                                                                                                                       Length 51;
                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                             49
                                                                                                                                                                                                                                                                                241
                                                                                                                                                                                                                                                                                                                  0
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AA173524
ID AA17
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AC AA17
AC AA17
AX
DT 09-N
XX
DT 09-N
XX
DE Huma
XX
KW Huma
KW Prot
KW Quar
XX
PN W020
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          therapy, and in vaccine production. (I) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of polymorphic polypeptides. For example, (I) may be used to treat disorders by rectifying mutations or deletions in a patient's genome that affect the activity of polypeptides by expressing inactive proteins or to supplement the patients own production of polypeptide. Additionally, (I) and its complementary sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and therefore which patients may be in need of restorative therapy. The polypeptides encoded by (I) may be used as antigens in the production of antibodies specific for polymorphic polypeptides. The antibodies may also be used to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic polymo
                                                                                              Human; single nucleotide polymorphism; SNP; genome; protein therapy; vaccine; probe; diagnostic assay; quantitation; restorative therapy; polymorphic; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide sequences (I), which contain single nucleotide polymorphisms (SNPs).

AAM53114 to AAM53329 represent peptides related to human polymorphic polynucleotide sequences. The sequences can be used in gene and protein therapy, and in vaccine production. (I) and the polypeptides encoded by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          protein therapy; vaccine; probe; diagnostic assay; detection;
quantitation; restorative therapy; polymorphic; ds.
                                                                                                                                                                                                       Human silent SNP containing nucleic acid SEQ:465.
                                                                                                                                                                                                                                                             09-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                   AAI73524 standard; DNA; 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 1; Page 2356; 2653pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2001-356160/37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    30-NOV-1999; 99US-0168138P.
29-NOV-2000; 2000US-00726173.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          polypeptides in samples
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Polymorphic nucleic acid sequences, useful in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30-NOV-2000; 2000WO-US032758
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (CURA-) CURAGEN CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  944 CCAGGCTGGAGTGCAATGGCCAAATCTCGGCTCACTGCAACCTCTGCCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CCAGGCTGGAGTGCAATGGTGTGATCTCGGCTCACTGCAACCTCCGCCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4.1%;
nilarity 89.8%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP; 12 A; 15 C; 16 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Leach
                                                                                                                                                                                                                                                                                                                                                                   ВÞ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; Score 41; DB;
; Pred. No. 2.1e.
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2.1e+02;
5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               genetic testing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                              ; gene therapy; detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     992
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WO200140521-A2 Homo sapiens

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RESULT 157
AAH89516/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CC AAMS3114 to AAMS329 represent peptides related to human polymorphic CC polynucleotide sequences. The sequences can be used in gene and protein CC therapy, and in vaccine production. (I) and the polymorphic may be used in the prevention, diagnosts and treatment of diseases CC associated with inappropriate expression of polymorphic polypeptides. For CC example, (I) may be used to treat disorders by rectifying mutations or CC deletions in a patient's genome that affect the activity of polypeptides by expressing inactive proteins or to supplement the patients own CC production of polypeptide. Additionally, (I) and its complementary CC sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and CC untibodies may be in need of restorative therapy. The CC antibodies specific for polymorphic polypeptides. The antibodies may also be used to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the presence of polymorphic colymorphic acids as a diagnostic agents for detecting the presence of polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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29-NOV-2000; 2000US-00726173:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Shimkets RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 51 BP; 10 A; 14 C; 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide sequences (I), which contain single nucleotide polymorphisms (SNPs).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; Page 196; 2653pp; English
                                                                                                                                                                                                                                                                                                                     Human; single nucleotide polymorphism; SNP; forensic test; aberrant protein expression;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Polymorphic nucleic acid sequences, useful in genetic testing
                           07-JAN-2000; 2000US-0174962P
                                                                                        05-JAN-2001; 2001WO-US000322.
                                                                                                                                                                                                                                                                 Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAH89516
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAH89516 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    polypeptides in
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                                                                                                                                                                                                          WO200151670-A2
                                                                                                                                                19-JUL-2001
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                                                                                                                                                                                                                                                                                                                                                                                                           coding sequence polymorphic site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            44;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Leach
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       samples
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          51
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Pred. No. 2.1e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                G; 14 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Db -,
. 2.1e+02;
5;
                                                                                                                                                                                                                                                                                                                                                                                                              SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                        paternity test;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 51;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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Matches
                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                     The present invention relates to human nucleic acids containing single nucleotide polymorphisms (SNPs). These can be used in forensic and paternity tests, and to aid in the treatment of diseases associated with aberrant protein expression, including cancer, amyloidosis, diabetes, Alzheimer's disease, Down's syndrome, oedema, lupus (SLE), vasculitis, glomerulonephritis, haemolytic anaemia, thrombocytopaenia, arthritis, meningitis, muscular disorders, dementia, nerrological diseases, tuberous sclerosis, male infertility, hypercalcaemia, blood pressure disorders, osteoporosis, pathogenic infections, hypercholesterolaemia, obesity or autoimmunity. The present sequence is a polymorphism-containing oligonucleotide fragment of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; Page 189; 475pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            polymorphisms, useful infection and diabetes.
                                                                                                                                                                                                         Sequence 51 BP; 11 A; 10 C; 20 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Shimkets
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Isolated human polynucleotides containing single nucleotide polymorphisms, useful for the treatment and diagnosis of e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     P-PSDB; AAM00399.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (CURA-) CURAGEN CORP.
                                                                                                                              Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2001-451871/48.
                              990 CCTCCCGGGCTCAAGCGATTCTCCTGTCTCAGCCTCCCAAGCAGCTGGG
  49
                                                                                                        44;
                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RA,
CCTCCCAGGTTCAAGCAATTCTCCTGCCTCAGCCTCCCAAGTAGCTGGG
                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Leach MD
                                                                                                                              4.1%;
                                                                                                      °.
                                                                                                                              Score 41;
Pred. No.
                                                                                                           Mismatches
                                                                                                                                                               DB 1;
                                                                                                                                    .1e+02;
                                                                                                                                                             Length
                                                                                                           Indels
                                                                                                                                                               51;
                                                         1038
                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cancer,
                                                                                                           Gaps
                                                                                                              0
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RESULT 158
AAH38408/c
14-AUG-2001
                   AAH38408;
                                      AAH38408 standard;
(first entry
                                       DNA;
                                       51
                                       뫔
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밁 Ś

Single nucleotide polymorphism; our; ourself, diabetes insipidus; cancer; SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; polycystic kidney disease; rheumatoid arthritis; multiple sclerosis; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis; Single nucleotide polymorphism; SNP; single nucleotide primer extension;

Human SNP flanking oligonucleotide SEQ ID 1204.

Homo sapiens.

WO200129262-A2

26-APR-2001

13-OCT-2000; 2000WO-US028436

15-OCT-1999; 99US-0160096P.

ORCHID BIOSCIENCES INC

Picoult-Newburg L,

WPI; 2001-290930/30

New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nuc acid sample. in a nucleic

Claim 1; Page 66; 83pp; English.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 159
AAH40504/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CC SNP flanking sequence, the SNPE primer is used as a genotyping primer.

CC The oligonucleotides are useful for genotyping a nucleic acid sample by performing a single-nucleotide primer extension reaction. The genotyping primer is used as a genotyping primer.

CC oligonucleotides are useful for determining the presence, absence or component to a sample of the genotyping nucleic acid samples, for e.g. to component to a sample of an individual or group of component to the genotypic traits suspected of being caused by one or more SNPs. Phenotypic traits include diseases e.g. component, diabetes insipidus, Lesch-Nyhan syndrome, muscular complex for susceptibility to multifactorial component is or may be genetic stickidney disease, confilated symptoms of or susceptibility to multifactorial confilammation, cancer, nervous system diseases and infection by pathogenic microorganism. The method is also useful in forensic investigations and component sequence represents a fragment of human component 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genotyping; agammaglobulinaemia; diabettes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis; inflammation; forensic investigation; paternity analysis; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide primer extension (SNPE) primers, and the sequences of regions flanking sites of single nucleotide polymorphisms SNPs. The present invention includes kits for determining the presence or absence of a SNP, using the oligonucleotides of the invention. The PCR primers are used to amplify a
                  New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nuc
                                                                                                                                                                                                                                                                                                      13-OCT-2000; 2000WO-US028436.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human SNP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             14-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAH40504 standard; DNA; 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
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                                                                                                  WPI; 2001-290930/30.
                                                                                                                                                                                                                                                       15-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                           WO200129262-A2
                                                                                                                                                                                                       (ORCH-) ORCHID BIOSCIENCES INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1024 TCCCAAGCAGCTGGGATTACGGGCACCTGCCACCACCCCGCTAATTTTT 1074
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TCCTRAGTAGCTGGGATTACAGGCACCTGCCACCACGCCCGGCTAACTTTT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        flanking
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP; 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         56; 83pp; English.
                                                                                                                                                                                                                                                         99US-0160096P
                                                                                                                                                  ŗ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          oligonucleotide SEQ ID 3300.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; 11 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4.1%;
                                                                                                                                                    3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17 G; 10 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 41;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 6,
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Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide primer extension (SNPE) primers, and the sequences of regions flanking controlled includes kits for determining the presence or absence of a SNP, using the coligonucleotides of the invention. The PCR primers are used to amplify a CC oligonucleotides of the invention. The PCR primers are used to amplify a CC oligonucleotides are useful for genotyping a nucleic acid sample by performing a single-nucleotide primer extension reaction. The CC oligonucleotides are useful for determining the presence, absence or cc identity of a SNP and for genotyping nucleic acid sample by clidantity of a SNP and for genotyping nucleic acid samples, for e.g. to casess by association analysis the genotype of an individual or group of clidatividuals, having a pathological phenotypic traits suspected of being caused by one or more SNPs. Phenotypic traits include diseases e.g. cagammaglobulinaemia, diabetes insipidus, Lesch-Nyhan syndrome, muscular cc dystrophy, familial hypercholesterolaemia, polycystic kidney disease, steegenesis imperfecta and acute intermittent porphyria. Phenotypic traits also include symptoms of or susceptibility to multifactorial cd disease of which a component is or may be genetic such as autoimmune cliseases, including, rheumatoid arthritis, multiple sclerosis, includes and include also useful in forensic investigations and conternity analysis. The present sequence represents a fragment of human content the airs of a sincle relation of human of the mineral content as a content of human of the present sequence represents a fragment of human content the airs of a sincle relation of the mineral content of the present sequence represents.
                                                                                                                                                                           Sequence 51
                                                                                                                                                                                                                                                                             flanking the site of a single nucleotide polymorphism
                                                                                                                                                                           BP;
                                                                                                                                                                           13
                                                                                                                                                                           A; 13 C; 15 G; 9 T;
4.1%;
Score 41;
Pred. No.
                                                                                                                                                                           0 U; 1 Other;
                                                                    51
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밁 Ś

Query Match Best Local S Matches 44

Similarity

44;

Conservative

Mismatches

0;

Gaps

0;

2.1e+02; 멂 ۲, 6,

Length Indels

ABL00045/c ID ABL000 RESULT 160 Human; single nucleotide polymorphism; SNP; polymorphism; cytostatic; immunosuppressive; antiinflammatory; neuroprotective; antimicrobial; autoimmune disease; inflammation; cancer; nervous system disease; Human silent 05-MAR-2002 ABL00045 standard; DNA; 51 infection; polymorphic protein; ds. (first entry) noncoding SNP oligonucleotide SEQ ID NO:36

Homo sapiens.

WO200138586-A2

22-NOV-2000; 2000WO-US032311

24-NOV-1999; 99US-0167383P

(CURA-) CURAGEN CORP.

RΑ, Leach

2001-355949/37.

Isolated human nucleic acids comprising one or more single nucleotide polymorphisms, useful for treating a subject suffering from a patholog e.g. autoimmune diseases, ascribed to the presence of a sequence polymorphism. pathology,

nucleic

absence or i

Claim 1;

Page 255;

674pp; English

ABL00010 to

ABL01104 represent human nucleic acid oligonucleotides

ABB56531

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RESULT 161
AAZ68649
ID AAZ686
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  and antibodies from the present invention can be used for treating a subject suffering from, at risk for, or suspected of, suffering from a pathology ascribed to the presence of a sequence polymorphism. The pathology may be autoimmune diseases, inflammation, cancer, diseases of the nervous system, and infection by pathogenic microorganisms. The SNPs are also useful for determining which forms of a characterised polymorphism are present in individuals. The antibodies may be used in the detection, quantitation and/or cellular or tissue localisation of a polymorphic protein (e.g., for use in measuring levels of the polymorphic protein within appropriate physiological samples)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              comprising one or more single nucleotide polymorphisms (SNPs). ABB56 to ABB56903 represent human peptides encoded by some of the SNP oligonucleotides. The sequences from the present invention can have
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 immunosuppressive, cytostatic, antiinflammatory, neuroprotective and antimicrobial activities. Nucleic acids, polypeptides, oligonucleotic
                                                                                                                                                                                                                                                                                                                                                                                                                                       Human genome, biallelic marker; high density disequilibrium map; genomic map; haplotype; phenotype; polymorphic base; genotyping; haplotyping; hybridisation; identification; characterisation; diagnosis; single nucleotide polymorphism; SNP; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 51 BP; 14 A; 13 C; 16 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                     21-APR-1998;
23-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human map-related biallelic marker SEQ ID NO:2999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAZ68649 standard;
                                                                                                                                                                                                                                   21-APR-1999;
                                                                                                                                                                                                                                                                  28-OCT-1999
                                                                                                                                                                                                                                                                                                WO9954500-A2
                                                                                                                                                                                                                                                                                                                                                              variation
                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10-SEP-2001
                                                                                             WPI; 2000-013267/01
                                             map of the human genome
                                                              Novel biallelic markers used to construct a high density disequilibrium
                                                                                                                                                        (GEST ) GENSET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  839 TCTGCCTGCCTCGGCCTCCCAAAGTGCTGGGATTACAGGCGTGAGCCAC 887
                                                                                                                         'n
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TCCTCCTGCCTTGGCCTCCCAAAGTTCTGGGATTATAGGCGTGAGCCAC 3
              Page 870;
                                                                                                                          Blumenfeld M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4.1%;
nilarity 89.8%;
Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                     98US-0082614P
98US-0109732P
                                                                                                                                                                                                                                   99WO-IB000822
                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers replace(24,T) /*tag= a
                                                                                                                                                                                                                                                                                                                               /standard_name= "single nucleotide polymorphism"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA;
              2745pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     47
                                                                                                                            Chumakov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
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Pred. No. 2.1e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 51;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
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RESULT 162
ADF77198
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Best Local S
Matches 43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAZ65654 to AAZ69578 represent human biallelic markers from the present invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AAZ69579 to AAZ77440 represent amplification primers for the biallelic markers. The biallelic markers of the invention primers to the stallelic markers. The biallelic markers of the invention have a variety of uses they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the compositions and methods of the invention can also be useful for the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            identification of the targets for the development of pharmaceutical agents and diagnostic methods, as well as the characterisation of the differential efficacious responses to and side effects from pharmaceutical agents acting on a disease as well as other treatment. N.B. The SEQ ID NOS 2852, 2913, 2974, 3035, 3096, 3157, 3227, 3297 and 3367, are not actually given a sequence in the Sequence Listing from the sequence of the se
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ss; single nucleotide polymorphism; KALPA; chromosome 13; membrane protein; SREBP; sterol response element binding protein transmembrane domain; TPR moutf; WD-40 respeat protein; aldo-keto reductase; AKR; domain; cholesterol-lowering compound;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 47 BP; 9 A; 16
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                                                                                                                                                                                                         Identifying a candidate cholesterol-lowering compound for treating cholesterol-related disorder by determining whether the test compounding the ability of the polypeptide to bind to the respective inhibits the ability of the polypeptide to bind to the respective inhibits.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    29-APR-2002; 2002US-0376510P
09-AUG-2002; 2002US-0402290P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cholesterol-related disorder; heart disease; coronary artery disease;
myocardial infarct; lipid-related metabolic disorder; obesity; diabetes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               KALPA SNP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     26-FEB-2004 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADF77198 standard;
                                                                                                                                                                                                                                                                                                                                                                           Benjamin S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         29-APR-2003; 2003WO-IB002376
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               13-NOV-2003.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           type
                                                                                                                                                                                  target polypeptide.
                                                                                                                                                                                                                                                                                                                          WPI; 2003-903719/82.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                 (CLIN-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ħ.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         839 TCTGCCTGCCTCGGCCTCCCAAAGTGCTGGGATTACAGGCGTGAGCC
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No. 2.1e+02;
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This sequence represents a fragment of the KALPA gene arou of a single nucleotide polymorphism (SNP). The KALPA gene chromosome 13 between markers D138265 and D138158. KALPA is

of the KALPA gene around the position

18

located on

D13S158. KALPA is a membrane element binding protein)

(sterol response

Disclosure; Fig 18; 196pp; English

protein involved

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RESULT 163
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ID AD1125
XX AD1125
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XX AD1125
XX BOT 22-APR
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KW breast
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Best Local S
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gene, such that the recommence to breast and ovarian cancerdeletion indicates a predisposition to breast and ovarian cancerdesent invention describes newly discovered deletion mutations in
                                                   This invention relates to a novel method for predicting a predisposition to cancer in a patient by detecting large deletions in the human tumour suppressor gene identified as the breast cancer susceptibility gene 1 (BRCA1). Specifically, it refers to deletions that result from the unequal crossover between a pair of repetitive Alu sequences in the BRCA1 gene, such that the recombined nucleotide sequence containing the deletion indicates a predisposition to breast and ovarian cancer. The
                                                                                                                                                                                                                                                                                                                 Predicting a predisposition to cancer in a patient comprising detecting deletion in the BRCA1 gene that results from the unequal crossover between a pair of repetitive sequences in the BRCA1 gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Scholl T,
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09-AUG-2002;
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breast cancer susceptibility gene 1; BRCA1; repetitive
ovarian cancer; junction sequence; recombination; mutar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human BRCA1
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                                                                                                                                                                                                                                                                                                                                                                                                                                  2004-062369/06
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2002US-0402430P.
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                                                                                                                                                                                                                                                                      English
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Best Local
                                                                                             The invention relates to a system for detecting gene expression, which comprises one or two isolated DNA molecules that detect expression of a gene, where the gene corresponds to any of 8143 oligonucleotides (ABZ00010-ABZ08152) each having 50 base pairs (bp). The system is useful for leukocyte expression profiling. It is particularly useful for diagnosing a disease, monitoring (rate of) progression of a disease, predicting therapeutic outcome, determining prognosis for a patient, predicting disease complications in an individual or monitoring respons to treatment in an individual. The diseases include cardiac allograft rejection, kidney allograft rejection, liver allograft rejection, atherosclerosis, congestive heart failure, systemic lupus erythematosus the program of the systemic lupus erythematosus the systemic lupus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               structure or biochemical function of BRCA1. Accordingly, it provides methods for detecting such mutants, as well as identifying and screening for cytostatic compounds useful for treating or preventing cancers associated with a BRCA1 genetic variant. This polynucleotide is a DNA fragment representing a junction sequence that arises as a result of a recombination event in human BRCA1 that causes the omission of exons 16 and 17, given in an exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New system for leukocyte expression profiling, diagnosing a disease, or monitoring (the rate of) progression of a disease, e.g. atherosclerosis or congestive heart failure, comprises diagnostic oligonucleotides.
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08-JUN-2001;
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atherosclerosis; congestive heart failure; systemic lupus erythematosus;
rheumatoid arthritis; osteoarthritis; cytomegalovirus; infection; probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-636525/68.
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Ly N, Woodward
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RESULT 165
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ABL59100/c
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                                                                                                                                                                                                                                      This is the nucleotide sequence for the PCR primer used in the amplification of the Alu repeat sequence, which is used to demonstrate the processes described in the invention. It involves the creation and use of circular yeast artificial chromosome (YAC) to selectively clone specific nucleic acids from a background of mixed nucleic acids by introducing the vector(s) into B. coli cells. They can be used to rapidly isolate human DNA where only a part of the sequence of DNA is known. Using the methods large fragments of DNA can be easily cloned and
                                                                                                                                                                                                                                                                                                                                                                            Preparation of yeast artificial chromosomes - by in vivo recombination using vector comprising yeast centromere, marker, yeast telomere and nucleic acid for recombination.
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  27-SEP-2002
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                                                                                                                                                                    Local
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slar yeast artificial chromosome; YAC; ss.
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llarity 100.0%;
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2e+02;
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Matches
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Best Local S
                                                                                                                                                                                                                               method, designated transformation-associated recombination, eliminates the need for an in vitro ligation step, and makes possible selective cloning of cDNAs for which only the 3'-sequence is known. The method is used for making a YAC. The method is also used for selective cloning of mammalian, specifically human, nucleic acid from a population, particularly radiation hybrids that contain only a small fragment of a human chromosome. The present sequence represents an Alu PCR primer. It was used for inter-Alu PCR, to produce ALu profiles of YACs produced using the method of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                The specification describes a method for making a yeast artificial chromosome (YAC) that includes an origin of replication (ori). The method comprises incorporating into yeast cells: a population of mammalian nucleic acid; and a vector that comprises a yeast centromere, selection marker, yeast telomere and a sequence that recombines with a region of the nucleic acid, so that in vivo recombination to a YAC occurs. This the nucleic acid, so that in vivo recombination to a YAC occurs.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Preparing yeast artificial chromosomes, useful e.g. for cloning specific human nucleic acid, comprises recombination in yeast cells between a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21-MAY-2002
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                                                                                                                                                                                          Sequence 40
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                                                                                                                                                                 Human sulphotransferase SULT1C1 gene polymorphic site,
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Cone polymorphism in such drug metabolising enzyme-encoding genes. The colymorphism is such drug metabolising enzyme-encoding genes. The colymorphism specific for a sequence selected from ABZ43217-ABZ50887 using a covariety of detection assays, including hybridisation assays, nucleic acid covariety of detection assays, including hybridisation assays, nucleic acid covariety of detection assays, including hybridisation assays, nucleic acid covariety of detection assays, including hybridisation assays, nucleic acid covariaty and processory, including hybridisation assays, nucleic acid covariaty and processory acid polymorphism data. Genetic polymorphism data, particularly that relating to single nucleotide polymorphism and ata, particularly that relating to single nucleotide covariations and human diseases, conditions, and responses to covary and sequence variations and human diseases. SNPs are particularly useful in the above respects as they are stable in populations, occur can all comparts and have lower mutation rates than other genome variations and penes encoding drug metabolising enzymes allows the customisation of cover the respecting sequences. The detection and analysis of polymorphisms covariate particular partients of covariations and the particular partient, but would also reduce the likelihood of adverse reactions, the thereby increasing safety. Covariate particular partient, but would also covariate the invention are also useful in the drug discovery and caparoval processes. For example, individuals could be selected for colinical trials only if their genetic profiles indicate that they are caparoval processes. For example, individuals could be selected for colonical trials only if their genetic profiles indicate that they are consisted and compositions of the invention may therefore lead to a an increase in the range of consisted with more detections, falsed drug trials, the time taken for a drug to be approved, the length of time patient seeds to take before finding an effective therapy in the col
                                                                                                                                         Matches
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02-MAY-2001;
27-AUG-2001;
                                                                                                                                                                                                                                                                                  Sequence 41
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    to methods and compositions for identifying individuals who have at least
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200252044-A2
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                                                                                                                                                                           Local
40
                                                                                                                                         40;
                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      s ABZ43217-ABZ50887 represent polymorphic sites within enzymes associated with drug metabolism. The invention
                                                                GCCTCGGCCTCCCAAAGTGCTGGGATTACAGGCGTGAGCC 885
                                                                                                                                                                                                                                                                                  BP; 7 A; 13 C; 14 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2000JP-00399443.
2001JP-00135256.
2001JP-00256862.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  즛
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          'standard_name= "Single nucleotide polymorphism (SNP)"
                                                                                                                                                                           100.0%;
                                                                                                                                                                                                              4.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Iida A,
                                                                                                                                 .0
                                                                                                                                                                               Score 40;
Pred. No.
                                                                                                                                         Mismatches
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                                                                                                                                                                       2e+02;
                                                                                                                                         0,
                                                                                                                                                                                                          Length 41
                                                                                                                                         Indels
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                                                                                                                                         Gaps
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27-DEC-2000; 2000JP-00399443
02-MAY-2001; 2001JP-00135256
27-AUG-2001; 2001JP-00256862
                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; drug metabolising enzyme; gene; drug metabolism; chromos polymorphic site; drug evaluation; drug screening; genotyping; genetic profiling; therapeutic customisation; adverse reaction;
                                                                                                                                                                                                                                                                           Nakamura Y,
                                                                                                                                                                                                                                                                                                                                                            WO200252044-A2
                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                          clinical trial; drug approval; single nucleotide polymorphism; SNP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human sulphotransferase SULTIC1 gene polymorphic site,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          26-JUN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABZ43598;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABZ43598 standard; DNA; 41 BP
                                                                                                                                                                                                                                                             WPI; 2002-583571/62
                                                                                                                                                                                                                                                                                        (RIKE ) RIKEN KK.
                                                                                                                                                                                                                                                                                                                                 27-DEC-2001; 2001WO-JP011592
                                                                                                                                                                                                                                                                                                                                                                                        variation
                                                                                                                                                                                                                                                                           Sekine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers replace(21,C)
                                                                                                                                                                                                                                                                                                                                                                         /standard_name= "Single nucleotide polymorphism (SNP)"
                                                                                                                                                                                                                                                                           Iida
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              #382
                                                                                                                                                                                                                                                                                                                                                                                                                                                chromosome
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Identifying individuals having a polymorphism, useful for determining the effectiveness or side effect of a drug or treatment protocol, comprises detecting at least one polymorphism in the drug metabolizing enzyme

Claim 23; Page 70; 2785pp; English.

sequences ABZ43217-ABZ50887 represent polymorphic sites within genes cc encoding enzymes associated with drug metabolism. The invention relates cc to methods and compositions for identifying individuals who have at least compositions in such drug metabolising enzyme-encoding genes. The cc primers specific for a sequence selected from ABZ43217-ABZ50887 using a cyariety of detection assays, including hybridisation assays, nucleic acid arrays and PCR-based methods. The invention also encompasses methods of cevaluating and screening drugs using genetic polymorphism data, particularly that relating to single nucleotide cc polymorphism data, particularly that relating to single nucleotide complymorphism data, particularly that relating to single nucleotide complymorphism data, particularly that relating to single nucleotide complymorphism and so useful as polymorphism markers for discovering genes to drugs. SNPs are also useful as polymorphism markers for discovering genes complys. SNPs are also useful as polymorphism markers for discovering genes comply to a particular particularly useful in the above respects as they are stable in populations, occur such as repeating sequences. The detection and analysis of polymorphisms complymorphism sequences and have lower mutation rates than other genome variations of drug therapies based upon the genetic profile of individual patients. This would not only take the guesswork out of selecting the drug with the greatest therapeutic effect for a particular patient, but would also createst therapeutic effect for a particular patient, but would also reduce the likelihood of adverse reactions, thereby increasing safety. Methods of the invention are also useful in the drug discovery and compositions of the invention are also useful in the drug discovery and compositions of the invention may therefore lead to a an increase in the range of the invention of the inventions. The methods, data and compositions of the invention of the inventions of the inventions of the inventions of the in

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RESULT 168

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Best Local S
Matches 40
                                                       Query Match
Best Local S
Matches 43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   possible drug targets and decreases in the number of adverse drug reactions, failed drug trials, the time taken for a drug to be approved, the length of time patients are on medication and the number of different medications a patient needs to take before finding an effective therapy
                                                                                                                                      The present invention relates to human nucleic acids containing single nucleotide polymorphisms (SNPs). These can be used in forensic and paternity tests, and to aid in the treatment of diseases associated with aberrant protein expression, including cancer, amyloidosis, diabetes, Alzheimer's disease, Down's syndrome, cedema, lupus (SLE), vasculitis, glomerulonephritis, haemolytic anaemia, thrombocytopaenia, arthritis, meningitis, muscular disorders, dementia, neurological diseases, tuberous sclerosis, male infertility, hypercalcaemia, blood pressure disorders, osteoporosis, pathogenic infections, hypercholesterolaemia, obesity or autoimmunity. The present sequence is a polymorphism-containing oligonucleotide fragment of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 07-JAN-2000; 2000US-0174962P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO200151670-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; single
forensic test;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human coding sequence polymorphic site SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAH89819;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAH89819 standard; DNA; 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 41 BP; 7 A; 13 C; 14 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Shimkets RA, Leach
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            05-JAN-2001; 2001WO-US000322
                                                                                                                                                                                                                                                                                                                        Claim 1; Page 277; 475pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19-JUL-2001
                                                                                                                                                                                                                                                                                                                                                                  Isolated human polynucleotides containing single nucleotide polymorphisms, useful for the treatment and diagnosis of e.g.
                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (CURA-) CURAGEN CORP
                                                        Local Similarity
mes 43; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    846 GCCTCGGCCTCCCAAAGTGCTGGGATTACAGGCGTGAGCC 885
                                                                                                                                                                                                                                                                                                                                                                                                                             2001-451871/48.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   40;
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                  GCCACCACACCCCGCTAATTTTTGTATTTTCATTAGAGGCGGGGTTTC 1099
                                                                                                                  50
   GCCACCACACCCAGCTAATTTTTGTATTTTAATAGAGACGGGGATTC
                                                                                                                                                                                                                                                                                                                                                   and diabetes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4.0%; ilarity 100.0%; Conservative
                                                                                                               B₽;
                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                nucleotide polymorphism; SNP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 aberrant protein expression;
                                                                                                                  14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          ₹
                                                                       4.0%;
89.6%;
                                                                                                                 12 C; 9 G; 15 T; 0 U; 0 Other;
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Pred. No.
                                                                       Score 40; I
Pred. No. 2
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                                                           Mismatches
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2e+02;
                                                     2.4e+02;
5;
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                                                                                                                                                                                                   neurological diseases, tuberous
                                                                                       ۳
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  paternity test; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Indels
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                                                                                     Length
                                                           Indels
   50
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                                                          0
                                                                                                                                                                                                                                                                                                                                                                      cancer,
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В Ś

0

28-AUG-2003 ACC84471;

(first entry)

ACC84471 standard;

DNA;

39 ВP

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RESULT 170
ACC84472
RESULT 171
ACC84471
ID ACC844
XX
AC ACC844
XX
DT 28-AUC
XX
                                                                                                                                                                                                                                                                                                                       Best Loca
Matches
                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to a neural thread protein (NTP) peptide referred to as cell death peptide. Thought to be cycostatic, antibacterial, immunosuppressive and antiinflammatory. It is useful for treating a condition in a patient requiring removal or destruction of cells, for treating a condition such as benign or malignant tumor, inflammatory disease, autoimmune disease and infectious disease. The peptide useful for treatment is derived from the amino acid sequence for a pancreatic thread protein. The peptide is conjugated, linked or bound to a molecule chosen from antibody or its fragment, antibody-like binding molecule, where the molecule has a higher affinity for binding to a tumor or other target than binding to other cells. Treatment using NTP peptides can remove benign tumors with less risk and fewer of the undesirable side effects of surgery. The present sequence is an NTP encoding sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19-JUL-2001;
19-JUL-2001;
16-NOV-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel neural thread protein peptide, referred as cell death peptide, useful for treating prostatic hyperplasia, psoriasis, eczema, dermatos atherosclerosis, cosmetic modification to skin, throat, mouth, muscle.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cytostatic; Antibacterial; Immunosuppressive; Antiinflammatory; neural thread protein; NTP; tumour; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NTP peptide encoding sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 39 BP; 6 A; 13 C; 12 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Averback
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19-JUL-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 19; 77pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   P-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (-OMYN)
                                                                                                                                                                                                                                                                                                                                               Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2003-247999/24.
                                                                                                                                                                                                                                                                            843 CCTGCCTCGGCCTCCCAAAGTGCTGGGATTACAGGCGTG
                                                                                                                                                                                                                                                                                                                          39;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       XOMYN
                                                                                                                                                                                                                                                                                                                                               Similarity
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                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2001US-0306150P.
2001US-0306161P.
2001US-0331477P.
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                                                                                                                                                                                                                                                                                                                                             100.0%;
                                                                                                                                                                                                                                                                                                                                                                     3.9%;
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                                                                                                                                                                                                                                                                                                                                               Score 39;
Pred. No.
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                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                        DB 1;
                                                                                                                                                                                                                                                                                                                                                  2.2e+02;
                                                                                                                                                                                                                                                                                                                               0
                                                                                                                                                                                                                                                                                                                                                                          Length 39;
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RESULT 172
ABA96813
ID ABA968
XX ABA968
XX ABA968
XY 30-APR
XX Human
XX Human;
KW Human;
KW Human;
KW immune
KW immune
XW immune
XW antiin
XX
PN WO2001
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        antibacterial, immunosuppressive and antiinflammatory. It is useful for treating a condition in a patient requiring removal or destruction of cells, for treating a condition such as benign or malignant tumor, inflammatory disease, autoimmune disease and infectious disease. The peptide useful for treatment is derived from the amino acid sequence for a pancreatic thread protein. The peptide is conjugated, linked or bound to a molecule chosen from antibody or its fragment, antibody-like binding molecule, where the molecule has a higher affinity for binding to a tumor or other target than binding to other cells. Treatment using NTP peptides can remove benign tumors with less risk and fewer of the undesirable side effects of surgery. The present sequence is an NTP encoding sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19-JUL-2001;
19-JUL-2001;
16-NOV-2001;
                                  Homo sapiens
                                                                  antiinflammatory;
                                                                                                                                                Human uteroglobin 9 probe,
                                                                                                                                                                                  30-APR-2002
                                                                                                                                                                                                                                                 ABA96813 standard; DNA; 41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to a neural thread protein (NTP) referred to as cell death peptide. Thought to be cytostatic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 19; 77pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel neural thread protein peptide, referred as cell death peptide, useful for treating prostatic hyperplasia, psoriasis, eczema, dermatos atherosclerosis, cosmetic modification to skin, throat, mouth, muscle.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19-JUL-2002; 2002WO-CA001105
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cytostatic; Antibacterial; Immunosuppressive; Antiinflammatory; neural thread protein; NTP; tumour; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (NYMO-) NYMOX CORP.
                                                             man; uteroglobin 9; recombinant production; malignant tumour; cancer;
lood disease; HIV infection; gene therapy; human immunodeficiency virus;
muune disorder; inflammatory condition; cytostatic; anti-HIV;
ntiinflammatory; immunomodulator; probe; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                l Similarity
39; Conserv
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                                                                                                                                                                                                                                                                                                                                                     CCTGCCTCAGCCTCCCAAGTAGCTGGGACCAAAGACATG 575
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                                                                                                                                                                                                                                                                                                                               CCTGCCTCAGCCTCCCAAGTAGCTGGGACCAAAGACATG 39
                                                                                                                                                                                                                                                                                                                                                                                              3.9%; So liarity 100.0%; F Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP; 10
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2001US-0306161P.
2001US-0331477P.
                                                                                                                                                                                (first entry)
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                                                                                                                                                SEQ ID NO:9.
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                                                                                                                                                                                                                                                                                                                                                                                                             Score 39;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             G; 6 T;
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                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                             0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                             DB 1; Le
2.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                             Length 39;
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RESULT 173
ABA96812
ABA96882
XX ABA968
XX ABA968
XX ABA968
XX ABA968
XX ABA968
XX Human
XX Human
XX Homoo s
XX immune
XX W2001
XX W2001
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to human uteroglobin 9 (AAM549078), nucleic acids encoding it (ABA96807), and a method for the recombinant production of uteroglobin 9. The protein has a molecular weight of 9 kD. The present invention additionally discloses an antagonist of uteroglobin 9 for therapeutic use, and an antibody which specifically binds to uteroglobin 9. Uteroglobin 9, and nucleotides which encode it may be used for treating a variety of diseases, such as malignant tumours, blood diseases, HIV (human immunodeficiency virus) infection, immune disorders and inflammatory conditions. The protein may also be used to screen for modulators of its activity or for peptide fingerprinting identification. The polynucleotide can be used as a primer for nucleic acid amplification reactions or as a probe for hybridisation reactions, or in producing gene chips or microarrays. Sequences ABA96812-ABA96813 represent human uteroglobin 9 probes used in an exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; uteroglobin 9; recombinant production; malignant tumour; cancer; blood disease; HIV infection; gene therapy; human immunodeficiency virus; immune disorder; inflammatory condition; cytostatic; anti-HIV;
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      мао у,
                                                                                                                                                                                              14-MAY-2001; 2001WO-CN000756
                                                                                                                                                                                                                                                            27-DEC-2001
                                                                                                                                                                                                                                                                                                                           WO200198337-A1
                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 antiinflammatory; immunomodulator; probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human uteroglobin 9 probe,
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                                                                    (SHAN-) SHANGHAI BIOWINDOW GENE DEV INC.
                                                                                                                                16-MAY-2000; 2000CN-00115717
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ilarity 95.1%;
Conservative
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Pred. No. 2.6e+02;
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RESULT 174
ABZ44526
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Best Local S
Matches 39
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 41 BP; 4 A; 10 C; 13 G; 14 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       and inflammatory conditions. The protein may also be used to screen for modulators of its activity or for peptide fingerprinting identification. The polynucleotide can be used as a primer for nucleic acid amplification reactions or as a probe for hybridisation reactions, or in producing gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 6; Page 19; 35pp; Chinese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human uteroglobin 9 and encoding polynucleotide, used in diagnosis and treatment of malignant tumors, hemopathy, human immunodeficiency virus infection, immunological diseases and inflammation.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          chips or microarrays. Sequences ABA96812-ABA96813 represent humanuteroglobin 9 probes used in an exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                         Human; drug metabolising enzyme; gene; drug metabolism; chromosome polymorphic site; drug evaluation; drug screening; genotyping; genetic profiling; therapeutic customisation; adverse reaction; clinical trial; drug approval; single nucleotide polymorphism; SNP,
                                                                                                     27-DEC-2000; 2000JP-00399443.
02-MAY-2001; 2001JP-00135256.
27-AUG-2001; 2001JP-00256862.
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                WPI; 2002-583571/62
                                            Nakamura Y,
                                                                                                                                                                     27-DEC-2001; 2001WO-JP011592
                                                                                                                                                                                                                                                                                               variation
                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human neuropathy target
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                                                                           RIKEN
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                                               Sekine
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                                                                                                                                                                                                                                                                             Location/Qualifiers replace(21,G) /*tag= a
                                                                                                                                                                                                                                                                /standard_name= "Single nucleotide polymorphism (SNP)"
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95.1%;
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Pred. No. 2
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Identifying individuals having a polymorphism, useful for determining the effectiveness or side effect of a drug or treatment protocol, comprises detecting at least one polymorphism in the drug metabolizing enzyme mucleic acid.
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2785pp; English

CC evaluating and screening drugs using genetic polymorphism data. Genetic CC polymorphism data, particularly that relating to single nucleotide CC polymorphisms (SNPs), may be used in studying the relationship between CC DNA sequence variations and human diseases. Conditions, and responses to CC drugs. SNPs are also useful as polymorphism markers for discovering genes that cause or exacerbate certain diseases. SNPs are particularly useful in the above respects as they are stable in populations, occur frequently, and have lower mutation rates than other genome variations of in genes encoding drug metabolising enzymes allows the customisation of CC drug therapies based upon the genetic profile of individual patients. CC This would not only take the guesswork out of selecting the drug with the CC greatest therapeutic effect for a particular patient, but would also CC reduce the likelihood of adverse reactions, thereby increasing safety. CC dinical trials only if their genetic profiles indicate that they are CC clinical trials only if their genetic profiles indicate that they are capable of responding to a particular drug ordrug class, and previously failed drug candidates could be revived if they were matched with more capable drug trials, the time taken for a drug to be approved, the length of time patients are on medication and the number of different medications a patient needs to take before finding an effective therapy tx one polymorphism in such drug metabolising enzyme-encoding genes. The polymorphisms may be identified in a nucleic acid sample using probes or primers specific for a sequence selected from ABZ4217-ABZ50887 using a variety of detection assays, including hybridisation assays, nucleic acid arrays and PCR-based methods. The invention also encompasses methods of Sequences ABZ43217-ABZ50887 represent polymorphic sites within genes encoding enzymes associated with drug metabolism. The invention relates to methods and compositions for identifying individuals who have at least Sequence 41 BP; 7 A; 16 C; 7 G; 11 T; 0 U; 0 Other;

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RESULT 175
ABZ50785
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Best Local :
                                                                           polymorphic
genetic prof
                                                                                                                 Human
                                                                                                                                   26-JUN-2003
                                                                                                                                                                          ABZ50785 standard; DNA;
                   variation
                                              Homo sapiens
                                                                                                                                                                                                                         676 CACTGCAACCTCTGCCTCCCGGGTTCAAGTTATTCTCCTGC 716
                                                                                                                neuropathy target esterase NTE gene polymorphic site,
                                                                                   drug metabolising enzyme; gene; drug metabolism; chromosome 19; rphic site; drug evaluation; drug screening; genotyping;
                                                                                                                                                                                                                                                                          Similarity
                                                                            prof
                                                                                                                                                                                                                                                                  Conservative
                                                                            lling; therapeutic customisation;
                                                                                                                                    (first entry)
                                                                drug approval;
         replace(21,G)
/*tag= a
                           Location/Qualifiers
                                                                                                                                                                                                                                                                          3.8%;
95.1%;
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                                                                                                                                                                           ВP
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Pred. No. 2
                                                                 single
                                                                                                                                                                                                                                                                   Mismatches
                                                                 nucleotide polymorphism;
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                                                                                                                                                                                                                                                                                     DB
                                                                            adverse reaction;
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                                                                   SNP; ds
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/standard_name= "Single nucleotide polymorphism (SNP)"

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                                                                                                                                                                                                                                                                                                       Sequences ABZ43217-ABZ50887 represent polymorphic sites within genes cencoding enzymes associated with drug metabolism. The invention relates to methods and compositions for identifying individuals who have at least cone polymorphism in such drug metabolising enzyme-encoding genes. The copymorphisms may be identified in a nucleic acid sample using probes or primers specific for a sequence selected from ABZ4217-ABZ50887 using a copymorphism data, particularly that relating to single nucleic acid evaluating and screening drugs using genetic polymorphism data. Genetic copymorphism data, particularly that relating to single nucleotide copymorphism data, genetic polymorphism data cause or exacerbate certain diseases. SNPs are particularly useful copymorphism strates than other genome variations and have lower mutation rates than other genome variations of that cause or exacerbate certain diseases. SNPs are particularly useful copymorphisms of genes encoding drug metabolising enzymes allows the customisation of corrections of the rappear dupymorphisms of polymorphisms of polymorphisms of genes encoding drug metabolising enzymes allows the customisation of corrections of adverse reactions, thereby increasing safety. Corrections of the invention are also useful in the drug discovery and approval processes. For example, individuals could be selected for capaboroval processes. For example, individuals could be selected for different copymorphisms of the patients are on methods, data and compositions of the inventions of the patients are on medication and the number of different medications, failed drug trials to take before fi
                                                         RESULT 176
                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                              Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Identifying individuals having a polymorphism, useful for determining the effectiveness or side effect of a drug or treatment protocol, comprises detecting at least one polymorphism in the drug metabolizing enzyme
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2002-583571/62
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02-MAY-2001; 2001JP-00135256
27-AUG-2001; 2001JP-00256862
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                                                                                                                                                                                                                                                                              16 C; 7 G; 11 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Iida
                                                                                                                                                                                                0;
                                                                                                                                                                                                                    Score 37.8;
Pred. No. 2.
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                                                                                                                                                                                                Mismatches
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                                                                                                                                                                                                                    .6e+02
                                                                                                                                                                                                                                   Length 41;
                                                                                                                                                                                                  Indels
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ADI12521 standard; DNA; 42

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RESULT 177
AAZ68006
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                                                                                                                                                                                                                                                                                                                        Query Match
Best Local S
Matches 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This invention relates to a novel method for predicting a predisposition to cancer in a patient by detecting large deletions in the human tumour suppressor gene identified as the breast cancer susceptibility gene 1 (BRCA1). Specifically, it refers to deletions that result from the unequal crossover between a pair of repetitive Alu sequences in the BRCA1 gene, such that the recombined nucleotide sequence containing the deletion indicates a predisposition to breast and ovarian cancer. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                present invention describes newly discovered deletion mutations that are believed to be deleterious and cause significant alterations in the structure or biochemical function of BRCA1. Accordingly, it provides methods for detecting such mutants, as well as identifying and screening for cytostatic compounds useful for treating or preventing cancers associated with a BRCA1 genetic variant. This polynucleotide is a DNA fragment representing a junction sequence that arises as a result of a fecombination event in human BRCA1 that causes the omission of exons 16
                                                                                                                                                                                                                                                                                                                                                                                           Sequence 42 BP; 8 A; 12 C; 12 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Predicting a predisposition to cancer in a patient comprising detecting deletion in the BRCA1 gene that results from the unequal crossover between a pair of repetitive sequences in the BRCA1 gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ds; cancer; human; tumour suppressor;
breast cancer susceptibility gene 1; BRCA1; repetitive Alu;
ovarian cancer; junction sequence; recombination; mutant.
Human genome; biallelic marker; high density disequilibrium map; genomic map; haplotype; phenotype; polymorphic base; genotyping;
                                                    Human map-related biallelic marker SEQ ID NO:2353.
                                                                                                                                                              AAZ68006 standard; DNA; 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 16; SEQ ID NO 1; 59pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           07-JUN-2002; 2002US-0387132P.
09-AUG-2002; 2002US-0402430P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               09-JUN-2003; 2003WO-US018098
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (MYRI-) MYRIAD GENETICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                   17, given in an exemplification of
                                                                                                                                                                                                                                                                        848 CTCGGCCTCCCAAAGTGCTGGGATTACAGGCGTGAGCCACC 888
                                                                                                                                                                                                                                                                                                                          1 Similarity
39; Conserv
                                                                                                                                                                                                                                                     N
                                                                                                                                                                                                                                                     CTCGGCCTCCCAAAGTGCTGGGATTACAGGTGTGAGCCATC 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hendrickson
                                                                                                                                                                                                                                                                                                                        3.8%;
nilarity 95.1%;
Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                       (first entry)
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                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                        Score 37.8; DB 1;
Pred. No. 2.6e+02;
0; Mismatches 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                   the invention.
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                                                                                                                                                                                                                                                                                                                            2; Indels
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                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    the BRCA1
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RESULT 178
ABZ43589/c
ID ABZ435
XX ABZ435
XC ABZ435
XX 26-JUN
XX Human
DE Human
XX Human
KW polymc
KW genet
KW genet
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AA265654 to AA269578 represent human biallelic markers from the present invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AA269579 to AA277440 represent amplification primers for the biallelic markers. The biallelic markers of the invention have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmaceutical agents and disgrastic methods, as well as the characterisation of the differential efficacious responses to and side effects from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                haplotyping; hybridisation; identification; characterisation; diagnosis; single nucleotide polymorphism; SNP; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel biallelic markers used to construct a high density disequilibrium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2000-013267/01
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21-APR-1998;
23-NOV-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   pharmaceutical agents acting on a disease as well as other treatment. N.B. The SEQ ID NOS 2852, 2913, 2974, 3035, 3096, 3157, 3227, 3297 and 3367, are not actually given a sequence in the Sequence Listing from the sequence of the sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO9954500-A2
                                                                                     Human; drug metabolising enzyme; gene; drug metabolism; chromos polymorphic site; drug evaluation; drug screening; genotyping; genetic profiling; therapeutic customisation; adverse reaction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 47
                                                                                                                                                                                                    Human cerebroside transferase CST gene polymorphic site,
                                                                                                                                                                                                                                                            26-JUN-2003
                                                                                                                                                                                                                                                                                                                                                                         ABZ43589 standard;
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     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1006 GATTCTCTGTCTCAGCCTCCCAAGCAGCTGGGATTACGGGCAC 1049
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             40;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Page 732; 2745pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Blumenfeld M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP; 9 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          98US-0082614P
98US-0109732P
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/standard_name= "single nucleotide polymorphism"
                                                             drug approval; single
                                                                                                                                                                                                                                                                                                                                                                         DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16 C; 11 G; 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        90.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 37.6;
Pred. No. 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                T; 0
                                                                   nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             .9e+02;
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o
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Other;
                                                                polymorphism;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                     chromosome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                   SNP;
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ct that cause of cadethate certain disease. Save all particularly certain control of the above respects as they are stable in populations, occur can the above respects as they are stable in populations of polymorphisms con such as repeating sequences. The detection and analysis of polymorphisms con in genes encoding drug metabolising enzymes allows the customisation of cingens encoding drug metabolising enzymes allows the customisation of conflict only take the guesswork out of selecting the drug with the conflict of the likelihood of adverse reactions, thereby increasing safety. Consider the likelihood of adverse reactions, thereby increasing safety. Confided the invention are also useful in the drug discovery and comportant processes. For example, individuals could be selected for comproval processes. For example, individuals could be selected for comproval processes. For example, individuals could be selected for compable of responding to a particular drug or drug class, and previously contained the populations. The methods, data and compositions of the invention may therefore lead to a na increase in the rumber of adverse drug creations, failed drug trials, the time taken for a drug to be approved, the length of time patients are on medication and the number of different confidence in the rumber of adverse drug to the length of time patients are on medication and the number of different confidence in the rumber of adverse of the length of time patients are on medication and the number of different confidence in the rumber of the particular confidence in the rumber of adverse drug to be approved.
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variation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    one polymorphisms may be identified in a nucleic acid sample using probes or primers specific for a sequence selected from ABZA3217-RAE50887 using a rvariety of detection assays, including hybridisation assays, nucleic acid arrays and PCR-based methods. The invention also encompasses methods of evaluating and screening drugs using genetic polymorphism data, particularly that relating to single nucleotide polymorphism (SNPs), may be used in studying the relationship between DNA sequence variations and human diseases, conditions, and responses to drugs. SNPs are also useful as polymorphism markers for discovering genes that cause or exacerbate certain diseases. SNPs are particularly useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequences ABZ43217-ABZ50887 represent polymorphic sites within genes encoding enzymes associated with drug metabolism. The invention rela
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Identifying individuals having a polymorphism, useful for determining the effectiveness or side effect of a drug or treatment protocol, comprises detecting at least one polymorphism in the drug metabolizing enzyme
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nakamura Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       27-DEC-2000; 2000JP-00399443
02-MAY-2001; 2001JP-00135256
27-AUG-2001; 2001JP-00256862
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     to methods and compositions for identifying individuals who have at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (RIKE ) RIKEN
medications a patient needs to take before finding an effective
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Page 70;
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/*tag= a
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문 S

41

Matches Query Match Best Local 8

Similarity

3.7%;

Conservative

<u>.</u>.

Pred. No. 3.1 0; Mismatches Score

: 36.2; No. 3

.1e+02 DB 1;

Length Indels

0;

Gaps

0

198 CATGITGGTCAGGCTGGTCTCGAACTCCCGACCTCAGATGA 238

Sequence 41 BP;

9 A; 11 C; 13 G; 8 T; 0 U; 0 Other;

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RESULT 179
ABZ45509
                                                                             variety of detection assays, including hybridisation assays, nucleic acid contrary and pCR-based methods. The invention also encompasses methods of evaluating and screening drugs using genetic polymorphism data, particularly that relating to single nucleotide polymorphism data, particularly that relating to single nucleotide polymorphisms (SNPs), may be used in studying the relationship between CC DNA sequence variations and human diseases, conditions, and responses to drugs. SNPs are also useful as polymorphism markers for discovering genes that cause or exacerbate certain diseases. SNPs are particularly useful in the above respects as they are stable in populations, occur frequently, and have lower mutation rates than other genome variations cuch as repeating sequences. The detection and analysis of polymorphisms in genes encoding drug metabolising enzymes allows the customisation of CC drug therapies based upon the genetic profile of individual patients. This would not only take the guesswork out of selecting the drug with the CC greatest therapeutic effect for a particular patient, but would also reduce the likelihood of adverse reactions, thereby increasing safety.

Methods of the invention are also useful in the drug discovery and control of course and the could be selected for
                                                                                                                                                                                                                                                                                                                                                                                                                                     to methods and compositions for identifying individuals who have at least one polymorphism in such drug metabolising enzyme-encoding genes. The polymorphisms may be identified in a nucleic acid sample using probes or primers specific for a sequence selected from ABZ43217-ABZ50887 using a variety of detection assays, including hybridisation assays
approval processes. For example, individuals could be selected for clinical trials only if their genetic profiles indicate that they are capable of responding to a particular drug or drug class, and previous failed drug candidates could be revived if they were matched with more failed drug candidates could be revived if they were matched with more
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequences ABZ43217-ABZ50887 represent polymorphic sites within g encoding enzymes associated with drug metabolism. The invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Identifying individuals having a polymorphism, useful for determining the effectiveness or side effect of a drug or treatment protocol, comprises detecting at least one polymorphism in the drug metabolizing enzyme
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02-MAY-2001; 2001JP-00135256.
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                                                                                             Sequences ABZ43217-ABZ50887 represent polymorphic sites within general encoding enzymes associated with drug metabolism. The invention relation methods and compositions for identifying individuals who have at
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one polymorphism in such drug metabolising enzyme-encoding genes. The polymorphisms may be identified in a nucleic acid sample using probes or primers specific for a sequence selected from ABZ43217-ABZ50887 using a variety of detection assays, including hybridisation assays, nucleic acid

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RESULT 181
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                                                                                     27-DEC-2000; 2000JP-00399443.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                    2001WO-JP011592
                                                                                                                                                                                                                                                                                                                                                replace(21,A)
/*tag= a
                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                                                                                                                                                                                                                                                       /standard_name= "Single nucleotide polymorphism (SNP)"
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92.7%;
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Pred. No. 3.1e+02;
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WP1; 2002-583571/62
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Identifying individuals having a polymorphism, useful for determining effectiveness or side effect of a drug or treatment protocol, comprise detecting at least one polymorphism in the drug metabolizing enzyme nucleic acid.

Claim 23; Page 197; 2785pp; English.

CC variety of detection assays, including hybridisation assays, nucleic acid carrays and pCR-based methods. The invention also encompasses methods of CC arrays and pCR-based methods. The invention also encompasses methods of CC polymorphism data, particularly that relating to single nucleotide polymorphism so (SNPs), may be used in studying the relationship between CC polymorphisms (SNPs), may be used in studying the relationship between CC DNA sequence variations and human diseases, conditions, and responses to that cause or exacerbate certain diseases, conditions, and responses to CC that cause or exacerbate certain diseases. SNPs are particularly useful cin the above respects as they are stable in populations, occur congresses encoding drug metabolising enzymes allows the customisation of CC greatest therapeutic effect for a particular partient polymorphisms CC in genes encoding drug metabolising enzymes allows the customisation of CC greatest therapeutic effect for a particular patient, but would also creduce the likelihood of adverse reactions, the drug with the CC greatest therapeutic effect for a particular patient, but would also ccur reduce the likelihood of adverse reactions, thereby increasing safety. CC enable of responding to a particular drug or drug class, and previously cfailed drug candidates could be revived if they were matched for capable of responding to a particular drug or drug class, and previously cfailed drug trials, the time taken for a drug to be approved, the length of time patients are on medication and the number of different medications a patient needs to take before finding an effective therapy or medications a patient needs to take before finding an effective therapy or consistent patient needs to take before finding an effective therapy or consistent provious thereby increases in the number of different medications a patient needs to take before finding an effective therapy or the patient provious therapy is a patient or take before finding an effective therapy or consistent p one polymorphism in such drug metabolising enzyme-encoding genes. The polymorphisms may be identified in a nucleic acid sample using probes primers specific for a sequence selected from ABZ43217-ABZ50887 using Sequences ABZ43217-ABZ50887 represent polymorphic sites within general encoding enzymes associated with drug metabolism. The invention related methods and compositions for identifying individuals who have at genes relates least õ

Sequence 41 BP; 9 A; 11 C; 13 G; 8 T; 0 U; 0 Other;

S Matches Query Match Local 198 CATGTTGGTCAGGCTGGTCTCGAACTCCCGACCTCAGATGA 238 38; Similarity 3.7**%**; 92.7**%**; Score 36.2; D Pred. No. 3.1e 0; Mismatches .1e+02 В Length 41; Indels 0, Gaps 0

RESULT 182 AAH91207/c

뭉

41

CATGTTGGCCAGGCTGGTCTCGAACTCCTGACCTCAGACGA 1

AAH91207 standard; DNA; 40

AAH91207;

09-OCT-2001 (first entry)

Human inflammatory bowel disease associated polymorphic site #282

Human; single inflammatory bowel disease; Crohn's disease; ulcerative colitis; nucleotide polymorphism; SNP; chromosome 19p13; paternity test; some 5q31-33; forensic test; gene therapy; ds.

Homo sapiens

misc_feature Location/Qualifiers

Nakamura

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Sekine A,

Iida

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Saito

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention describes a method for detecting the presence polymorphisms associated with inflammatory bowel diseases such as ulcerative colitis and Crohn's disease. The methods can be used to d the presence of genetic polymorphisms associated with inflammatory b disease and correlating their occurrence with disease states. They m
27-DEC-2000; 2000JP-00399443
                       27-DEC-2001; 2001WO-JP011592
                                                                                                                                     Key
                                                                                                                                                            Homo
                                                                                                                                                                                  Human; drug metabolising enzyme; gene; drug metabolism; chromosome polymorphic site; drug evaluation; drug screening; genotyping; genetic profiling; therapeutic customisation; adverge reaction; clinical trial; drug approval; single nucleotide polymorphism; SNP,
                                                                                                                                                                                                                                               Human NDUFS1
                                                                                                                                                                                                                                                                         26-JUN-2003
                                                                                                                                                                                                                                                                                                 ABZ50133;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Testing for the presence of polymorphisms associated with inflammatory bowel disease, using a hybridization assay.
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10-APR-2000; 2000US-0196046P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          polymorphic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        in this way for phenotypic correlations, forensics, paternitying, medicine and genetic analysis. The present sequence is a morphic site described in the exemplification of the invention
                                                                                                                                                            sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hudson TJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP; 13
                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                               gene
                                                                                                                                                                                                                                                                       (first entry)
                                                                                                            replace(21,T)
/*tag= a
                                                                                                                                  Location/Qualifiers
                                                                                            /standard_name= "Single nucleotide polymorphism (SNP)"
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                                                                                                                                                                                                                                               polymorphic site, #6915.
                                                                                                                                                                                                                                                                                                                        DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; 7 C; 12 G; 7 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                              3.6%;
92.5%;
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RESULT 184 ABZ44123/c

ABZ44123

standard; DNA; 41 BP

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40

1045 GGCACCTGCCACACCCCCGCTAATTTTTGTATTTTCA 1083

GGCACATGCCACCACCCGGCTAATTTTTGTATTTTCA

Best Local Similarity
Matches 37; Conserv

Conservative

0;

Mismatches

0;

Gaps

0

3.6%; 94.9%;

Score 35.8; Pred. No. 3

.3e+02

Length Indels

Query Match

26-JUN-2003

(first entry)

Human NDUFS1 gene polymorphic site, #907.

Human; drug metabolising enzyme; gene; drug metabolism; chromosome polymorphic site; drug evaluation; drug screening; genotyping;

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Sequences ABZ43217-ABZ50887 represent polymorphic sites within genes cencoding enzymes associated with drug metabolism. The invention relates to methods and compositions for identifying individuals who have at least cone polymorphism in such drug metabolising enzyme-encoding genes. The polymorphisms may be identified in a nucleic acid sample using probes or primers specific for a sequence selected from ABZ43217-ABZ50887 using a carrays and pCR-based methods. The invention also encompasses methods of evaluating and screening drugs using genetic polymorphism data, particularly that relating to single nuclected colymorphism data, particularly that relating to single nucleotide polymorphisms (SNPs), may be used in studying the relationship between CR polymorphisms (SNPs), may be used in studying the relationship between CR polymorphisms are also useful as polymorphism markers for discovering genes to that cause or exacerbate certain diseases, SNPs are particularly useful cin the above respects as they are stable in populations, cocur frequently, and have lower mutation rates than other genome variations come such as repeating sequences. The detection and analysis of polymorphisms come in genes encoding drug metabolising enzymes allows the customisation of CR drug therapies based upon the genetic profile of individual patients. CR mis would not only take the guesswork out of selecting the drug with the greatest therapies based upon the genetic profile of individual patients. CR mis would not only take the guesswork out of selecting the drug with the greatest therapies based upon the genetic profile of individual patients.

CR methods of the invention are also useful in the drug discovery and approval processes. For example, individuals could be selected for cindal patient populations. The methods data and compositions of the invention may therefore lead to a an increase in the range of the inventions of the methods. Adata and compositions of the inventions of the methods of adverse drug to be approved.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Identifying individuals having a polymorphism, useful for determining effectiveness or side effect of a drug or treatment protocol, comprise detecting at least one polymorphism in the drug metabolizing enzyme
Sequence 41 BP; 12 A; 7 C; 12 G; 10 T; 0 U; 0 Other;
                                                reactions, failed drug trials, the time taken for a drug to be approved, the length of time patients are on medication and the number of different medications a patient needs to take before finding an effective therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim
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27-AUG-2001; 2001JP-00256862.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 23; Page 206; 2785pp; English.
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CC sequences ABZ43217-ABZ50887 represent polymorphic sites within genes conding enzymes associated with drug metabolism. The invention relates cone polymorphism in such drug metabolising enzyme-encoding genes. The polymorphism shape identified in a nucleic acid sample using probes or complymorphism in such drug metabolising enzyme-encoding genes. The polymorphism shape identified in a nucleic acid sample using probes or complymorphism data, particularly their relation assays, nucleic acid carays and pcR-based methods. The invention also encompasses methods of cevaluating and screening drugs using genetic polymorphism data. Genetic polymorphism data, particularly that relating to single nucleotide complymorphism solutions and human diseases, conditions, and responses to drugs. SNPs are also useful as polymorphism markers for discovering genes conditions and human diseases. SNPs are particularly useful compasses or exacerbate certain diseases. SNPs are particularly useful in the above respects as they are stable in populations, occur frequently, and have lower mutation rates than other genome variations such human diseases. SNPs are particularly useful cause or exacerbate certain diseases. SNPs are particularly useful in the above respects as they are stable in populations, occur frequently, and have lower mutation rates than other genome variations genes encoding drug metabolising enzymes allows the customisation of drug therapeutic effect for a particular patient, but would also creduce the likelihood of adverse reactions, thereby increasing safety. Cc methods of the invention are also useful in the drug discovery and aproval processes. For example, individuals could be selected for capatroyriate patient populations. The methods, data and compositions of the lumpth of time patients are on medication and the number of different medications a patient needs to take before finding an effective therapy of different medications and the number of different medications and the number of different medications and the nu
                                       Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        27-DEC-2000; 2000JP-00399443
02-MAY-2001; 2001JP-00135256
27-AUG-2001; 2001JP-00256862
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clinical trial; drug approval; single nucleotide polymorphism; SNP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 23; Page 79; 2785pp; English
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3.6%;
nilarity 94.9%;
Conservative
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Score 35.8; DB 1;
Pred. No. 3.3e+02;
0; Mismatches 2;
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                                                                              G; 10 T; 0 U; 0 Other;
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                                           Length 41;
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Best Loc Matches

Local

l Similarity 37; Conser

2;

Indels

0

Gaps

0

RESULT 186
AAV19045/c
ID AAV190
XX
AC AAV190
XX
DT 28-JUL

AAV19045 standard; DNA; 40

28-JUL-1998 AAV19045;

(first entry)

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RESULT 185
AAT97407/c
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                                                          Matches
                                                                                      Query Match
                                                                                                                                                               A method has been developed for detecting the presence of a target site (TS), of at least one nucleotide (nt) in a nucleic acid (NA). The method comprises: (a) forming an oligonucleotide (ON), consisting of two fluorophores (F1, F2) each covalently linked to separate nt, bound to TS; and (b) detecting fluorescence energy transfer (FET) between F1 and F2 when ON is released from TS. The present sequence represents a synthetic polynucleotide used in an example of the present invention. The method is used to diagnose hereditary and other diseases; to determine infectious agents; in tissue typing for histocompatibility; in forensic identification and paternity testing, and in monitoring the genetic make up of plants and animals. Specifically it is used to detect single nt polymorphisms. The method provides inexpensive, simple, accurate and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic oligomer D18S8 Allele G from WO9722719 Example 2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAT97407;
                                                                                                                                                                                                                                                                                                                                                                                                                     Detecting target site in nucleic acid by forming a fluorophore-labelled oligonucleotide at the site - and detecting fluorescent energy following denaturation, used e.g. to detect inherited diseases, in tissue typing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Detection; target site; nucleic acid; fluorophore; labelled; fluorescent; inherited disease; tissue typing; PCR; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                          etc
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Kwok P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18-DEC-1995;
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                                                                                                                  Sequence 40 BP; 11 A; 7 C; 15
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                                                                          Local
                   675 TCACTGCAACCTCTGCCTCCCGGGTTCAAGTTATTCTCCT
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40
                                                                         Similarity
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 TCACTGCAAGCTCTGCCTCCCGGGTTCAAGCAATTCTCCT 1
                                                                                                                                                       nucleic
                                                            Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    95US-0008743P
                                                                                                                                                       acid analyses
                                                                          92.5%;
                                                             0;
                                                                          Score 35.2;
Pred. No. 3.
                                                                                                                        G; 7
                                                            Mismatches
                                                                                                                        T; 0 U; 0 Other;
                                                                              .4e+02
                                                                                            DB
                                                                                            1; Length
                                                               Indels
                                                                 0
                                                                 Gaps
                                                                 0,
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RESULT 187
ABL59101/c
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                                                                                                                                                                                                                                                                                                                                                                     Matches
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                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                      This is the nucleotide sequence for the PCR primer used in the amplification of the Alu repeat sequence, which is used to demonstrate the processes described in the invention. It involves the creation and use of circular yeast artificial chromosome (YAC) to selectively clone specific nucleic acids from a background of mixed nucleic acids by introducing the vector(s) into E. coli cells. They can be used to rapidly isolate human DNA where only a part of the sequence of DNA is known. Using the methods large fragments of DNA can be easily cloned and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Alu
          09-JUL-1996;
                                  14-APR-1998;
                                                                                   US6391642-B1
                                                                                                                                   transformation-associated recombination; PCR; primer; ss.
                                                                                                                                                  Yeast artificial chromosome; YAC; inter-Alu PCR;
                                                                                                                                                                          Nucleotide
                                                                                                                                                                                                   27-SEP-2002
                                                                                                                                                                                                                                                    ABL59101 standard; DNA; 40
                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 1; Page 45; 117pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  nucleic acid for recombination.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Preparation of yeast artificial chromosomes - by in vivo recombination using vector comprising yeast centromere, marker, yeast telomere and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1998-110234/10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Resnick MA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              09-JUL-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     09-JUL-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15-JAN-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO9801573-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Saccharomyces
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PCR; primer; amplification; Alu repeat sequence; vector; circular yeast artificial chromosome; YAC; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (USSH ) US DEPT HEALTH & HUMAN SERVICES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PCR
                                                                                                                                                                                                                                                                                                                                           987
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           vector comprising yeast centromere,
                                                                                                                                                                                                                                                                                                                    40
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                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                  CTGCCTCCCGGGCTCAAGCGATTCTCCTGTCTCAGCCTCC 1026
                                                                                                                                                                                                                                                                                                                  CCGCCTCCGGGTTCAAGCGATTCTCCTGCCTCAGCCTCC
                                                                                                                                                                        sequence of an Alu PCR primer
                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                     BP; 9 A; 8 C; 19 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Larionov
                                                                                                                                                                                                 (first entry)
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          96WO-US011478
                                   98US-00060023
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                                                                                                                                                                                                                                                                                                                                                                                3.6%;
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                                                                                                                                                                                                                                                                                                                                                                                Score 35.2;
Pred. No. 3.
                                                                                                                                                                                                                                                                                                                                                                    Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                 .4e+02;
                                                                                                                                                                                                                                                                                                                                                                                            DB 1; Length 40;
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                                                                                                                                                                                                                                                                                                                                                                    Gaps
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method, designated transformation-associated recombination, eliminates the need for an in vitro ligation step, and makes possible selective cloning of cDNAs for which only the 3'-sequence is known. The method is used for making a YAC. The method is also used for selective cloning of mammalian, specifically human, nucleic acid from a population, particularly radiation hybrids that contain only a small fragment of a human chromosome. The present sequence represents an Alu PCR primer. It was used for inter-Alu PCR, to produce ALu profiles of YACs produced
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      chromosome (YAC) that includes an origin of replication (ori). The metl comprises incorporating into yeast cells: a population of mammalian nucleic acid, and a vector that comprises a yeast centromere, selection marker, yeast telomere and a sequence that recombines with a region of the property of the control of the control of the property of the p
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Preparing yeast artificial chromosomes, useful e.g. for cloning specific human nucleic acid, comprises recombination in yeast cells between a nucleic acid and a yeast vector.
                                                                                              using the method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The specification describes a method for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 1; Col 27; 50pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2002-498777/53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (USSH ) US DEPT HEALTH & HUMAN SERVICES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Larionov
                                                                                                        of
f
                                                                                              the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Kouprina NY,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 making a yeast artificial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Perkins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         臣
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뭉 Ş Matches Query Match Local 987 CTGCCTCCCGGGCTCAAGCGATTCTCCTGTCTCAGCCTCC 1026 40 1 Similarity 37; Conserv CCGCCTCCCGGGTTCAAGCGATTCTCCTGCCTCAGCCTCC Conservative 3.6%; 0, Score 35.2; DB 1; Pred. No. 3.4e+02; Mismatches Length 40; Indels 0, Gaps 0

Sequence 40 BP; 9 A; 8 C; 19 G; 4 T; 0 U; 0 Other;

RESULT 188
AAH49727/c
ID AAH497
XX AAH497
XX AAH497
XX LIB Human;
XW Human;
XW Human;
XW Human;
XW Human
XX Human
XX Human
XX Human
XX IB Homo 8
XX IB Homo 9
XX Human; DNA repair mismatch protein 11; cancer; haemopathy; HIV infection; immunological disease; inflammation; gene therapy; probe; ss. Human DNA mismatch repair protein 11 coding sequence probe 25-SEP-2001 AAH49727 standard; (first entry) DNA; 41

WO200147988-A1

05-JUL-2001

18-DEC-2000; 2000WO-CN000627

23-DEC-1999; 99CN-00125733

(UYFU-) UNIV FUDAN. (SHAN-) SHANGHAI BI BIO DOOR GENE TECHNOLOGY LTD

Mao Y, Xie

WPI; 2001-425639/45

DNA mismatch repair protein 11 and encoded polynucleotide, diagnosis and treatment of malignant tumor, hemopathy, HIV immunological diseases and various inflammation. applicable infection, ä

Example

36pp;

Chinese.

The present invention provides the protein and coding sequences of human DNA mismatch repair protein 11. The sequences are useful in the treatment of cancer, haemopathy, HIV infection, immunological diseases and inflammation. The present sequence is a probe for the coding sequence of

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RESULT 189
ABL40963/c
ID ABL409
XX ABL409
XX ABL409
XX Transc
XX Transc
XX Transc
XX Transc
XX Homo s
XX W02002
XX W20002
XX W02002
XX W01; 2
XX W
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Matches 37
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Best Local Similarity
Matches 37; Conserv
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                                                                                                                                                                                                                                 The invention relates to a novel human transcription regulation factor ZFM1 isomer 19.47 and polynucleotides encoding the protein. The protein can be expressed by standard recombinant methodology. The protein and encoding polynucleotides are used in diagnosis and treatment of malignar tumours, haemopathy, HIV infection, immunological diseases and inflammation. The present sequence represents the human transcription regulation factor ZFM1 isomer 19.47 cDNA specific probe
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human transcription regulation factor ZFM1 isomer 19.47 and encoding polynucleotide, used in diagnosis and treatment of malignant tumors, hemopathy, human immunodeficiency virus infection, immunological diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Transcription regulation factor ZFM1 isomer 19.47; human; ZFM1; transcription regulation factor; cytostatic; haemostatic; viruc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14-MAR-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                            Example 6;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   inflammation.
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                                               708
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  41
                                                                                            l Similarity
37; Conserv
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                                                                                                                                                                                           41
                              TTCTCCTGCCCCAGCCTCCTGAGTAGCTGGGACTACAGGC
     TTCTCCTGCCTCAACCTCCCGAGTAGCTGGGACTACAGGC 2
                                                                                                                                                                                                                                                                                                                                                                                                                            Page 22; 38pp;
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                                                                                                                                                                                      B₽;
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                                                                                            Conservative
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                                                                                                                   3.6%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Chinese.
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Pred.
                                                                                               0
                                                                                            Score 35.2; DI Pred. No. 3.5e. 0; Mismatches
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No. 3
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                                                                                                                                            DB 1;
                                                                                                                                                                                           0 Other;
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                                                                                                                                            Length 41;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cDNA specific probe
                                                                                                 Indels
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                                                                                               0;
                                                                                                                                                                                                                                                                                                              of malignant
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                                                                                               Gaps
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03-MAR-2003 ABZ20666; ABZ20666

standard;

DNA;

41

Human; G protein subunit 9.02; cancer; constipation; cardiac asthma; colic; psychic disease; probe; morphinic analgesic acute poisoning; ss.

diarrhoea; cough;

G protein subunit 9-02 coding sequence probe

Homo sapiens

밁 δ

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ABL40964/c
ID ABL409
RESULT 191
ABZ20666/c
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                                                                                                                                                                                                                                                                                                   The invention relates to a novel human transcription regulation factor ZFM1 isomer 19.47 and polynucleotides encoding the protein. The protein can be expressed by standard recombinant methodology. The protein and encoding polynucleotides are used in diagnosis and treatment of malignan tumours, haemopathy, HIV infection, immunological diseases and inflammation. The present sequence represents the human transcription regulation factor ZFM1 isomer 19.47 cDNA specific probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human transcription regulation factor ZFM1 isomer 19.47 and encoding polynucleotide, used in diagnosis and treatment of malignant tumors, hemopathy, human immunodeficiency virus infection, immunological dise and inflammation.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        25-JUN-2001; 2001WO-CN001050
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Transcription
                                                                                                                                                                                                                                                             Sequence 41 BP; 9 A; 9 C; 15 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 6; Page 22; 38pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2002-258025/30.
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                                                                                                                                                                                                 Local
                                                                                                                                 708
                                                                                      41
                                                                                                                                                                           l Similarity
37; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Xie
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                                                                                                                                     TTCTCCTGCCCCAGCCTCCTGAGTAGCTGGGACTACAGGC
                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
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92.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        factor; cytostatic; haemostatic;
                                                                                                                                                                             0
                                                                                                                                                                                                 Score 35.2;
Pred. No. 3
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                                                                                                                                                                               Mismatches
                                                                                                                                                                                                   .5e+02
                                                                                                                                                                                                                       DB 1;
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                                                                                                                                                                                  Indels
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                                                                                                                                                                                  Gaps
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RESULT 192
ABQ77547
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                                                                                                                                                                                                                                                                                                                            Mao
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       G protein subunit 9.02. The sequences can be used in the treatment of cancers, coughs, cardiac asthma, diarrhoea, constipation, colic, psychic disease and morphinic analgesic acute poisoning. The present sequence is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The
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                              Example 6; Page 19
                                                                                                                                       New polypeptide-human red blood cell cytoplasmic treating anaerobic cerebral disease, respiratory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       23-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23-AUG-2000; 2000CN-00119732.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  respiratory adynacardiant; probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; red blood cell cytoplasmic protein 15.29; erythrocyte; recombinant production; gene therapy; cerebral anoxia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human red blood cell cytoplasmic protein 15.29 probe,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-OCT-2002
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                                                                                                         intestinal
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                                                                                                                                                                                                                                                                                                                                                                                                (BODE-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     probe used to isolate the coding sequence of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local
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                                                                                                                                                                                                                                                    2002-472206/51.
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37; Conserv
                                                                                                                                                                                                                                                                                                                            Xie
                                                                                                                                                                                                                                                                                                                                                                                                BODE GENE DEV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            6; Page 22 (Disclosure);
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SHANGHAI
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                                                                                                     palsy, and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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                              (Disclosure); 32pp;
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                                                                                                         anemia.
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Pred. No. 3.
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                                                                                                                                       protein 15.29 for
adynamia, arrhythmia,
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                                                                                                                                                                                                                                                                                  polynucleotide, and a method for preparing the polypeptide by DNA recombination. The application of the polypeptide is in treating arrhythmia and diabetes. Also disclosed are the antagonist against the polypeptide and its therapeutic action, and the application of the polynucleotide. The current sequence represents a human protein 10.01
                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to a human protein designated 10.01, cont 
Phe-His aminolyase active site. Also disclosed are the encoding
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                                                                                                                                                                                                                                                              related
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                              655 TGCAGTGGCGCAATCTTGGCTCACTGCAACCTCTGCCTCC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     7; Page 21 (disclosure); 33pp; Chinese.
                                                                                                                                                                                                                                                            probe sequence
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TGCAGTGGCGCAATCTTGGTTCACTGCAACCCCCCCCCTCC
                                                                                                                                                                                                          BP; 6 A;
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                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 aminolyase active site; arrhythmia; diabetes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10.01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DEV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3.6%;
92.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     related probe
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                                                                                                                           3.6%;
                                                                                                                                                                                                          16 C; 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              41
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                                                                                                     Score 35.2; DB 1;
Pred. No. 3.5e+02;
0; Mismatches 3
                                                                                                                                                                                                          G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SHANGHAI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 35.2;
Pred. No. 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          .5e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ريا
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40
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         e site and diabetes.
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RESULT 195
AAQ27391
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                                                                                                                                                                                                                                                                              Query Match
Best Local (
                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                           The present invention discloses a polypeptide-guanosine triphosphatase activator protein-IO.01. The invention also discloses the method for curing several diseases, such as squamobasal cell carcinoma of skin, osteosarcoma, leukemia and teratoma by using said polypeptide. The present sequence represents a probe for guanosine triphosphate activations.
                                                                                                                                                                                                                                                                                                                                                                                                                                  A polypeptide-guanosine triphosphatase activator protein -10.01 polynucleotide for coding this polypeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        мао Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20-NOV-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CN1380320-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            carcinoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Guanosine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Probe #1 for guanosine triphosphate activator 10.01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14-JUL-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ACC00156 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10-APR-2001; 2001CN-00105912
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10-APR-2001; 2001CN-00105912
                                                                                                                                                                     AAQ27391 standard;
                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                            Example 7; Page 23; 33pp; Chinese.
                                                                                                                   25-MAR-2003
27-JAN-1993
                               WO9213101-A1
                                                    Synthetic
                                                                         Polymerase
                                                                                             Inter-Alu specific primer PDJ33.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (SHAN-)
                                                                                                                                                                                                                                                655 TGCAGTGGCGCAATCTTGGCTCACTGCAACCTCTGCCTCC 694
                                                                                                                                                                                                                           \vdash
                                                                                                                                                                                                                                                                                                                                    sequence
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                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                           TGCAGTGGCACAGTCTCGGCTCACTGCAACCTCTGCCTCC 40
                                                                                                                                                                                                                                                                                                               41 BP; 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          triphosphatase activator 10.01; squamobasal cell; of skin; osteosarcoma; leukemia; teratoma: prohe-
                                                                         chain
                                                                                                                                                                                                                                                                      Conservative
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                                                                                                                 (revised)
(first en
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                                                                         reaction;
                                                                                                                                                                                                                                                                                                               A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             osteosarcoma; leukemia; teratoma; probe;
                                                                                                                                                                                                                                                                               3.6%;
                                                                                                                                                                       DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  entry)
                                                                                                                   entry)
                                                                                                                                                                                                                                                                                                               16 C; 10 G; 9 T; 0 U; 0 Other;
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                                                                         PCR; repetitive element;
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Pred. No. 3.5e
0; Mismatches
                                                                                                                                                                                                                                                                                .5e+02;
                                                                                                                                                                                                                                                                                            DB 1;
                                                                                                                                                                                                                                                                                         Length 41;
                                                                                                                                                                                                                                                                                                                                               polypeptide. The triphosphate activator
                                                                                                                                                                                                                                                                       Indels
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RESULT 196
ABQ83633
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Best Local S
Matches 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Detection of genetic variation by 2-D electrophoresis of fragments - and hybridisation with labelled probes, carried out on fragments consisting of inter-repeat sequences generated by PCR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Primer PDJ33 is one of several primers which are preferred for use in amplifying inter-Alu regions of DNA. The amplified fragments are then subjected to 2-D electrophoresis on the basis of length and differences in base sequence. The resulting separation pattern is transferred to a filter for screening with a probe. The method can be used to detect genetic variation. See AAQ27339-Q27404 and AAQ33141-Q33144. (Updated on
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 6; Page 6; 31pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1992-284683/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 35 BP; 8 A; 10 C; 11 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-JAN-1991;
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                                                                                                                                                                                                           24-APR-2002.
                                                                                                                                                                                                                                CN1345805-A.
                                                                                                                                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                                                                                                      endocrinopathy;
                                                                                                                                                                                                                                                                                               Human; mPer3-10.01;
                                                                                                                                                                                                                                                                                                                     Human mPer3-10.01 probe 1 SEQ ID NO:8.
                                                                                                                                                                                                                                                                                                                                           26-JAN-2003
                                                                                                                                                                                                                                                                                                                                                                  ABQ83633;
                                                                                                                                                                                                                                                                                                                                                                                      ABQ83633 standard; DNA; 41 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25-MAR-2003 to correct PN field.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (INGE-) INGENY BV
                                                                                               WPI; 2002-539321/58
                                                                                                                   Mao Y,
                                                                                                                                                               26-SEP-2000;
                                                                                                                                                                                    26-SEP-2000;
                                                                                                                                         (SHAN-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Sim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   852
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                                                                                                                                         SHANGHAI BIOWINDOW GENE
                                                                                                                                                                                                                                                                                                                                                                                                                                                            GCCTCCCAAAGTGCTGGGATTACAGGCGTGAGCCA 886
                                                                                                                                                                                                                                                                                                                                                                                                                                              GCCTCCCAAAGTGCTGGGATTACAGGCGTGAGCCA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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G
                                                                                                                                                               2000CN-00125425.
                                                                                                                                                                                     2000CN-00125425
                                                                                                                                                                                                                                                                                                                                            (first entry)
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                                                                                                                                                                                                                                                                                      .01; vegetative nervous dysfunction; psychic disease; growth development disturbance disease; tumour; prob
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Vijg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 35; DB; Pred. No. 3.2
                                                                                                                                            VED
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3.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                        tumour; probe;
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The present invention describes human mPer3-10.01 a method for producing (I) using DNA recombination used in the treatment of several diseases, such as

DNA recombination diseases, such as

technology. (I) car vegetative nervous Also described hnology. (I) can

1 is 1 be

(I).

Novel polypeptide-human mPer 3-10.01 and polypeptide.

polynucleotide

for encoding

Example 6; Page 20 (Disclosure); 33pp; Chinese

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RESULT 197
ABQ83634
ID ABQ836
AC ABG836
AC ABG86
A
                                                                           RESULT 198
ABL52955/c
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Best Local S
Matches 37
                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local 9
                                                                                                                                                                                                                                                                                                      Best Local Similarity Matches 37; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention describes human mPer3-10.01 (I). Also described is a method for producing (I) using DNA recombination technology. (I) can be used in the treatment of several diseases, such as vegetative nervous dysfunction, psychic disease, endocrinopathy, growth development disturbance disease and tumours. The present sequence represents a probe for (I), which is used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         dysfunction, psychic disease, endocrinopathy, growth development disturbance disease and tumours. The present sequence represents a probe for (I), which is used in an example from the present invention
ABL52955;
                                                 ABL52955
                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 41 BP; 5 A; 10 C; 14 G; 12 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 6; Page 20 (Disclosure); 33pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel polypeptide-human mPer 3-10.01 and polynucleotide for encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2002-539321/58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              26-SEP-2000; 2000CN-00125425
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; mPer3-10.01; vegetative nervous dysfunction; psychic disease; endocrinopathy; growth development disturbance disease; tumour; probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human mPer3-10.01
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37; Conserv
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                                                    standard;
                                                                                                                                                                                                                        TGGAGTTTCTCCATGTTGGTCAGGCTGGTCTCGAACTCCCG 227
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TGGGGTTTCACCATGTTGGGCAGGCTGGTCTCGAACTCCTG 41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      regaerriciccarerreercaeecreercreaacreece 227
                                                                                                                                                                                           TGGGGTTTCACCATGTTGGGCAGGCTGGTCTCGAACTCCTG
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                                                    DNA;
                                                                                                                                                                                                                                                                                                                             3.5%;
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                                                    41
                                                    BP.
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                                                                                                                                                                                                                                                                                                 Score 34.6; DB 1;
Pred. No. 3.7e+02;
0; Mismatches 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DEV INC
                                                                                                                                                                                                                                                                                                                                                             DB 1;
                                                                                                                                                                                                                                                                                                      4; Indels
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                                                                                                                                                                                                                                                                                                                                                       Length 41;
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RESULT 199
ABZ49715
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                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mao Y,
                                                                                                                  Key
variation
                                                                                                                                                                            Human; drug metabolising enzyme; gene; drug metabolism; chromosome polymorphic site; drug evaluation; drug screening; genotyping; genetic profiling; therapeutic customisation; adverse reaction; clinical trial; drug approval; single nucleotide polymorphism; SNP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention relates to serine proteinase 10 (AAM48453). Serine proteinase 10 and its coding sequence can be used for treating diseases such as cancer and HIV infection. The present sequence is a probe, which was used in an example from the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Serine proteinase 10; enzyme; cancer; HIV infection; anti-HIV; cytostatic; probe; ss.
27-DEC-2000; 2000JP-00399443
                      27-DEC-2001; 2001WO-JP011592
                                              04-JUL-2002.
                                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                                                       Human sulphotransferase TPST2 gene polymorphic site, #6497
                                                                                                                                                                                                                                                                26-JUN-2003
                                                                                                                                                                                                                                                                                       ABZ49715;
                                                                                                                                                                                                                                                                                                           ABZ49715 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 41 BP; 9 A; 14 C; 13 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Polypeptide-serine proteinase 10 and polynucleotide encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2002-196715/26.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     31-MAY-2000; 2000CN-00116278
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           31-MAY-2000; 2000CN-00116278
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                                                                      WO200252044-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    12-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (BODE-) BODE GENE DEV CO LTD SHANGHAI.
                                                                                                                                                                                                                                                                                                                                                                                             637 CTGTCACCCAGGCTGGAGTGCAGTGGCGCAATCTTGGCTCA 677
                                                                                                                                                                                                                                                                                                                                                                      41
                                                                                                                                                                                                                                                                                                                                                                                                                    37;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Xie Y;
                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 6; Page 19 (Disclosure); 32pp; Chinese
                                                                                                                                                                                                                                                                                                                                                                       CTGTCGCCCAGGCTGGAGTGCAGTGGTGCCATCTCGGCTCA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                (first entry)
                                                                                                      Location/Qualifiers replace(21,A) /*tag= a
                                                                                            /standard_name= "Single nucleotide polymorphism
                                                                                                                                                                                                                                                                                                                                                                                                                               3.5%;
                                                                                                                                                                                                                                                                                                             41
                                                                                                                                                                                                                                                                                                             ВÞ
                                                                                                                                                                                                                                                                                                                                                                                                                    Score 34.6; I
Pred. No. 3.7e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                     0
                                                                                                                                                                                                                                                                                                                                                                                                                                3.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1; Length 41;
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                                                                                                                                                                                                                                                                                                                                                                                                                     0,
                                                                                                                                                                              SNP;
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                                                                                             " (SNP)
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RESULT 200
ABZ43958
ID ABZ4395
XX
AC ABZ439
XX
DT 26-JUN
XX
DE Human
XX
KW Human;
KW polymc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequences ABZ43217-ABZ50887 represent polymorphic sites within genes encoding enzymes associated with drug metabolism. The invention relates to methods and compositions for identifying individuals who have at least cone polymorphism in such drug metabolism; enzyme-encoding genes. The cone polymorphisms may be identified in a nucleic acid sample using probes or primers specific for a sequence selected from ABZ43217-ABZ50887 sucleic acid carrays and PCR-based methods. The invention also encompasses methods of cevaluating and screening drugs using genetic polymorphism data, particularly that relating to single nuclectide polymorphism data particularly that relating to single nuclectide complymorphism data are also useful as polymorphism markers for discovering genes in the above respects as they are stable in populations, and responses to crugs. SNPs are also useful as polymorphism markers for discovering genes in the above respects as they are stable in populations, occur frequently, and have lower mutation rates than other genome variations in genes encoding drug metabolising enzymes allows the customisation of crugs with the genetic profile of individual patients. The detection and analysis of polymorphisms cour reduce the likelihood of adverse reactions, thereby increasing safety. Methods of the invention are also useful in the drug discovery and experience in the invention are also useful in the drug discovery and approval processes. For example, individuals could be selected for capable of responding to a particular patient, but would also creduce the likelihood of adverse reactions, thereby increasing safety. Creduce the likelihood of adverse reactions the drug discovery and approval processes. For example, individuals could be selected for capable of responding to a particular drug or drug class, and previously failed drug candidates could be reviewed if they were matched with more capable of the invention may therefore lead to a na increase in the trange of the inventions of the inventions. The methods, d
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                                                                                                                                                                                                                                                                                                                                                       S
                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local S
Matches 37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             effectiveness or side detecting at least one nucleic acid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 23; Page 196; 2785pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Identifying individuals having a polymorphism, useful for determining the effectiveness or side effect of a drug or treatment protocol, comprises detecting at least one polymorphism in the drug metabolizing enzyme
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nakamura Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             02-MAY-2001;
27-AUG-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 41 BP; 6
    Human; drug
polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  reactions, failed drug trials, the time taken for a drug to be approved, the length of time patients are on medication and the number of different medications a patient needs to take before finding an effective therapy
                                                                Human glutathione-S-transferase MGST2 gene polymorphic site, #742.
                                                                                                              26-JUN-2003
                                                                                                                                                                                                      ABZ43958 standard;
                                                                                                                                                                                                                                                                                                                                                         969 CTCGGCTCACTGCAACCTCTGCCTCCCGGGCTCAAGCGATT 1009
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                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                CTCGGCTCACTGCAACCTCCGCCTCCCGGGTTCAAGCAGTT 41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sekine
  metabolising enzyme; gene; drug metabolism; chromosome site; drug evaluation; drug screening; genotyping;
                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
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2001JP-00256862.
                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           P.
                                                                                                                                                                                                      DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                        3.5%;
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                                                                                                                                                                                                      41 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                          Score 34.6;
Pred. No. 3.
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                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1; Length 41;
                                                                                                                                                                                                                                                                                                                                                                                                        Indels
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CC to methods and compositions for identifying interviewed to one polymorphism in such drug metabolising enzyme-encoding genes. The compositions pecific for a sequence selected from ABZ4217-ABZ50887 using a cC variety of detection assays, including hybridisation assays, nucleic acid carrays and PCR-based methods. The invention also encompasses methods of CC evaluating and screening drugs using genetic polymorphism data. Genetic polymorphisms (SNPs), may be used in studying the relationship between CC polymorphisms (SNPs), may be used in studying the relationship between CC polymorphisms (SNPs), may be used in studying the relationship between CC that cause or exacerbate certain diseases, conditions, and responses to CC drugs. SNPs are also useful as polymorphism markers for discovering genes that cause or exacerbate certain diseases. SNPs are particularly useful CC in the above respects as they are stable in populations, occur in the above respects as they are stable in populations, occur in genes encoding drug metabolising enzymes allows the customisation of CC drug therapies based upon the genetic profile of individual patients. CC mis would not only take the guesswork out of selecting the drug with the CC greatest therapeutic effect for a particular patient, but would also reduce the likelihood of adverse reactions, thereby increasing safety. CC capable of responding to a particular drug or drug class, and previously failed drug candidates could be revived in the drug discovery and comportate patient populations. The methods, data and compositions of the invention may therefore lead to a an increase in the range of the inventions a patients are on medication and the number of different medications and the number of different medicati
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       27-DEC-2000; 2000JP-00399443.
02-MAY-2001; 2001JP-00135256.
27-AUG-2001; 2001JP-00256862.
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clinical trial; drug approval; single nucleotide polymorphism; SNP; ds.
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detecting at least
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  27-DEC-2001; 2001WO-JP011592
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2002-583571/62.
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/*tag= a
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Query Match Best Local S Matches 37

Similarity

3.5%; 15 C; 10

37;

Conservative

0,

Mismatches

Pred.

No. 3.7e+02

Score 34.6; DB 1; G; 10 T; 0 U; 0 Other;

Length 41; Indels

0

Gaps

0

Sequence 41

BP;

6 A;

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RESULT 201
ABZ4950
ID ABZ4950
XX ABZ4950
XX ABZ4950
XX ABZ4950
XX Human
XX Human
XX Human
XX Key
FT Vari
FT Vari
FT Vari
FT Vari
FT T Vari
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PR 27-
PR 27
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                Sequences ABZ43217-ABZ50887 represent polymorphic sites within genes
CC encoding enzymes associated with drug metabolism. The invention relates
CC to methods and compositions for identifying individuals who have at least
CC one polymorphism in such drug metabolising enzyme-encoding genes. The
CC primers specific for a sequence selected from ABZ43217-ABZ50887 using a
CC variety of detection assays, including hybridisation assays, nucleic acid
CC arrays and PCR-based methods. The invention also encompasses methods of
CC evaluating and screening drugs using genetic polymorphism data, particularly that relating to single nucleotide
CC polymorphisms (SNPs), may be used in studying the relationship between
CC DNA sequence variations and human diseases, conditions, and responses to
CC drugs. SNPs are also useful as polymorphism markers for discovering genes
CC that cause or exacerbate certain diseases. SNPs are particularly useful
CC frequently, and have lower mutation rates than other genome variations
CC such as repeating sequences. The detection and analysis of polymorphisms
CC drug therapies based upon the genetic profile of individual patients.
CC This would not only take the genetic profile of individual patients.
CC greatest therapeutic effect for a particular patient, but would also
CC methods of the invention are also useful in the drug with the
CC methods of the invention are also useful in the drug with the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Identifying individuals having a polymorphism, useful for determining effectiveness or side effect of a drug or treatment protocol, comprise detecting at least one polymorphism in the drug metabolizing enzyme
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nakamura
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   27-DEC-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; drug metabolising
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2001JP-00135256.
2001JP-00256862.
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invention are also useful in the

discovery

Sequences ABZ43217-ABZ50887 represent polymorphic sites within genes encoding enzymes associated with drug metabolism. The invention relation methods and compositions for identifying individuals who have at one polymorphism in such drug metabolishing enzyme-encoding genes. The polymorphisms may be identified in a nucleic acid sample using probes primers specific for a sequence selected from ABZ43217-ABZ50887 using

have at learnes. The

relates least

probes or

Identifying individuals having a polymorphism, useful for determining the effectiveness or side effect of a drug or treatment protocol, comprises detecting at least one polymorphism in the drug metabolizing enzyme nucleic acid.

23; Page

184;

2785pp; English.

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RESULT 202
RABZ49230
ID ABZ492
XX Human;
XX Human;
XX Human;
XX ABZ492
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       27-DEC-2000;
02-MAY-2001;
27-AUG-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       approval processes. For example, individuals could be selected for clinical trials only if their genetic profiles indicate that they are capable of responding to a particular drug or drug class, and previously failed drug candidates could be revived if they were matched with more appropriate patient populations. The methods, data and compositions of the invention may therefore lead to a an increase in the range of possible drug targets and decreases in the number of adverse drug reactions, failed drug trials, the time taken for a drug to be approved, the length of time patients are on medication and the number of different medications a patient needs to take before finding an effective therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; drug metabolising enzyme; gene; drug metabolism; poly drug evaluation; drug screening; genotyping; genetic profilitherapeutic customisation; adverse reaction; clinical trial;
                                                                                                                                                                                                                                                                                                                                                                                                                                     Nakamura
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         04-JUL-2002.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (RIKE ) RIKEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    27-DEC-2001; 2001WO-JP011592
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
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2001JP-00135256
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              standard_name= "Single nucleotide polymorphism (SNP)"
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Pred. No. 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            site,
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Nakamura Y, Sekine A,

Iida A,

Saito

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABZ49236
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                variety of detection assays, including hybridisation assays, nucleic actic arrays and PCR-based methods. The invention also encompasses methods of evaluating and screening drugs using genetic polymorphism data. Genetic polymorphism data, particularly that relating to single nucleotide polymorphisms (SNPs), may be used in studying the relationship between DNA sequence variations and human diseases, conditions, and responses to drugs. SNPs are also useful as polymorphism markers for discovering genes
                                                          27-DEC-2000;
02-MAY-2001;
27-AUG-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 41
                                                                                                                                                                                                                                                                                                                                                                                                                  Human; drug metabolising enzyme; gene; drug metabolism; polymor drug evaluation; drug screening; genotyping; genetic profiling; therapeutic customisation; adverse reaction; clinical trial;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human aldehyde dehydrogenase ALDH6Al gene polymorphic site, #6019.
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                                                                                                                                     27-DEC-2001; 2001WO-JP011592
                                                                                                                                                                               04-JUL-2002
                                                                                                                                                                                                                  WO200252044-A2
                                                                                                                                                                                                                                                                                               variation
                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
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                  (RIKE)
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                                                                                                                                                                                                                                                                                                                                                                                                 approval;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             992 TCCCGGGCTCAAGCGATTCTCCCTGTCTCAGCCCTCCCAAGCA 1032
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       37;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TCCCGGGTTCAAGCGATTCTCCTGCCTCAGCCTCCCGAGTA 41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    detection assays, including hybridisation assays, nucleic acid PCR-based methods. The invention also encompasses methods of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP; 6 A; 16 C; 9 G;
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                                                                             2000JP-00399443
2001JP-00135256
                                                                                                                                                                                                                                                                                                                                                                                                 single nucleotide polymorphism;
                                                                                                                                                                                                                                                                          Location/Qualifiers replace(21,C)
/*tag= a
                                                                                                                                                                                                                                                          /standard_name= "Single nucleotide polymorphism (SNP)
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No. 3
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                                                                                                                                                                                                                                                                                                                                                                                                      SNP; ds.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              polymorphic site;
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198 CATGTTGGTCAGGCTGGTCTCGAACTCCGACCTCAGATGA 238

Query Match Best Local Matches

37;

Conservative

0; Score Pred.

Mismatches

4.

Indels Length

0

Gaps

0

41;

34.6; No. 3. No.

.7e+02; В

Similarity

3.5%; 10 C;

Sequence 41 BP; 6

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13 G; 12 T;

0 U;

0 Other;

medications a patient needs to take before finding an effective therapy

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RESULT 204

ABZ43562 standard; DNA; 41 BP

Human 26-JUN-2003

sulphotransferase TPST2 gene polymorphic site,

(first entry)

Human; drug metabolising enzyme; gene; drug metabolism; chromosome polymorphic site; drug evaluation; drug screening; genotyping; genetic profiling; therapeutic customisation; adverse reaction; clinical trial; drug approval; single nucleotide polymorphism; SNP;

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CC evaluating and screening drugs using genetic polymorphism data. Genetic CC polymorphism data, particularly that relating to single nucleotide CC polymorphism data, particularly that relating to single nucleotide CC polymorphisms (SNPs), may be used in studying the relationship between CC DNA sequence variations and human diseases, conditions, and responses to CC drugs. SNPs are also useful as polymorphism markers for discovering genes that cause or exacerbate certain diseases. SNPs are particularly useful in the above respects as they are stable in populations, occur frequently, and have lower mutation rates than other genome variations such as repeating sequences. The detection and analysis of polymorphisms CC in genes encoding drug metabolising enzymes allows the customisation of CC drug therapeutic effect for a particular patient, but would also reduce the likelihood of adverse reactions, thereby increasing safety. CC reduce the likelihood of adverse reactions, thereby increasing safety. CC dapproval processes. For example, individuals could be selecting they are CC capable of responding to a particular drug or drug class, and previously capable of responding to a particular drug or drug class, and previously failed drug candidates could be revived if they were matched with more cc appropriate patient populations. The methods, data and compositions of the invention may therefore lead to a nincrease in the rumber of adverse drug creations, failed drug trials, the time taken for a drug to be approved, the length of time patients are on medication and the number of different medications and the number of different the length of time patients are on medication and an effective theraps.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               one polymorphism in such drug metabolising enzyme-encoding genes. The polymorphisms may be identified in a nucleic acid sample using probes primers specific for a sequence selected from ABZ43217-ABZ50887 using variety of detection assays, including hybridisation assays, nucleic a arrays and PCR-based methods. The invention also encompasses methods of the compasses of the compas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 encoding enzymes
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             represent
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metabolism. The invention rela
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Location/Qualifiers

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CC variety of detection assays including hybridisation assays, nucleic acid CC variety of detection assays, including hybridisation assays, nucleic acid CC evaluating and screening drugs using genetic polymorphism data. Genetic CC polymorphism data, particularly that relating to single nucleotide CC polymorphism (analysis), may be used in studying the relationship between CC drugs. SNPs are also useful as polymorphism markers for discovering genes that cause or exacerbate certain diseases, conditions, and responses to CC in the above respects as they are stable in populations, occur crequently, and have lower mutation rates than other genome variations in genes encoding drug metabolishing enzymes allows the customisation of CC includes based upon the genetic profile of individual patients. CC This would not only take the guesswork out of selecting the drug with the greatest therapeutic effect for a particular patient, but would also comproved processes. For example, individuals could be selected for clinical trials only if their genetic profile of individual patients. CC daparoval processes. For example, individuals could be selected for clinical trials only if their genetic profiles indicate that they are capable of responding to a particular drug or drug class, and previously failed drug candidates could be revived if they were matched with more appropriate patient populations. The methods, data and compositions of the invention may therefore lead to a an increase in the range of compositions a patient needs to take before finding an effective therapy was medications a patient needs to take before finding an effective therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         sequences ABZ43217-ABZ50887 represent polymorphic sites within genes encoding enzymes associated with drug metabolism. The invention relates to methods and compositions for identifying individuals who have at least one polymorphism in such drug metabolising enzyme-encoding genes. The polymorphisms may be identified in a nucleic acid sample using probes or primers specific for a sequence selected from ABZ43217-ABZ50887 using a variety of detection assays, including hybridication association.
                                                                                                                                                                                             Sequence
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02-MAY-2001; 2001JP-00135256
27-AUG-2001; 2001JP-00256862
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                                                                                              Similarity
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CTCGGCTCACTGCAACCTCTGCCTCCCGGGCTCAAGCGATT 1009
                                                                                                                                                                                             B₽;
                                                                Conservative
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The invention relates to an isolated nucleic acid or alternate splice covariant comprising a nucleotide sequence containing at least one of the single nucleotide polymorphisms given in the specification, a nucleotide complement of them. The genes are ADAM19 (a disintegrin and metalloprotease 19, also known as gene 845), NRG2 (neuroregulin 2, also known as gene 847), endophilin 1 (also known as gene 874), endophilin 2 (also known as gene 874), endophilin 2 (also known as gene 874), endophilin 2 (also known as gene 874), also included care a vector comprising the isolated nucleic acid (or alternate splice variant), a host cell containing the vector, an isolated polypeptide composition typel motif 2, also known as gene 962). Also included care a vector comprising the isolated nucleic acid (or alternate splice variant), an encophilin 1 (alternate splice variant), an encophilin 2 (also known as gene 962). Also included care avector composition to composition the vector, an isolated polypeptide composition of composition to alternate splice variant), an exception of antibody fragment that binds to the polypeptide, and a carrier, complice variant, vector, polypeptide or antibody, and a carrier, composition of component to detecting a disintegrin and mucleic acid or alternate splice variant, antibody or antibody fragment, and at least one component to detect the hybridisation of the variant or the binding of the antibody to an ADAM gene amino acid sequence), a kit for detecting an interactor gene amino acid sequence), a kit for detecting an interactor gene amino acid sequence), a kit for detecting an ADAM or interactor gene amino acid sequence), a kit for detecting an ADAM or interactor gene amino acid sequence), a kit contained to detect the indirance and sequence (comprising the contained and acid sequence), a kit contained to the interactor gene amino acid sequence), a kit contained to the interactor gene amino acid sequence) of an orthologue of a human subject, determining an ADAM or interactor gene, treating an AD
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Allen K;
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a disintegrin and metalloprotease with thrombospondin typel motif 2;
asthma; atopy; obesity; inflammatory bowel disease; respiratory disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New isolated nucleic acid or alternate splice variant, useful for diagnosing and treating a disintegrin and metalloprotease (ADAM) or interactor gene-associated disorder, e.g. asthma, atopy, obesity or inflammatory bowel disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Endophilin 1; Endophilin 2; NRG2; ADAMTS2;
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Best Local
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                                                         Edmonds M,
Tsuchihashi
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADL64137 standard;
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(PERR/)
(POWE/)
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                                                                                                                                                                                   RAMANATHAN C S.
SWANSON B.
TSUCHIHASHI Z.
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                                                                                                                                                                                                                                                                                                             HUI L.
PERRONE M.
                                                                                                                                                                                                                                                                                                                                                                    EDMONDS M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              41
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                                                                                             Hui
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                                                                   Zerba
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3.5%;
                                                                                                Perrone M,
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Pred. No. 3.
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                                                                                                Powell JR,
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                                                                                                   Ramanathan CS,
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                                                                                                   Swanson
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New nucleic acid comprising a single nucleotide polymorphism at a specific location, useful in paternity testing, genetic analysis or diagnosing, preventing or treating cardiovascular diseases e.g. angioedema or angina pectoris.
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SEQ ID 376pp; English.

CC (C1S), the alanyl aminopeptidase protein (ANDED), the meprin A, beta CC protein (MEPIB), the aminopeptidase protein (XPR), the meprin A, beta CC kallkrein protein (KLKI), the membrane bound aminopeptidase P protein (C (XPR)), the tissue (C (XPR)), the tissue (C (XPR)). The nucleic acid comprises at least one polymorphic position, CC (Including the alleles, reference alleles and alternate alleles of the CC single nucleotide polymorphisms, listed in the specification. The CC sequence of the gene. The polymorphic position residing in a coding consition resides within the untranslated composition resides within the untranslated region or an intronic region of the gene. The polymorphic position residing in a coding consition resides within the untranslated region or an intronic region of the gene. Constructing haplotypes using the nucleic acids above further comprises using the haplotypes to identify an individual for the presence of a disease phenotype, and correlating the presence of the disease constructing haplotypes to identify an individual for the presence of anilysis. The nucleic acids and polyperides can be used in diagnosing, correlating eracitovascular disease, primers and probes are comprised with the haplotype. The nucleic acids primers and probes are comprised mailysis. The nucleic acids and polypeptides can be used in diagnosing, correlating cardiovascular disease, e.g. andiocedema, angina pectoris, hypersension, heart failure, myocardial infarction, aneurysm, stroke, embolism, thrombosis, coronary artery disease or cancer. The cost of the invention.

CC arteriosclerosis, hypersensitivity reactions during haemodialysis. Cost the invention. The invention relates to an isolated nucleic acid (I) derived human gene encoding a protein, such as the C1, S subcomponent human gene encoding (C1S), the alanyl ar cid (I) derived from a S subcomponent protein

Sequence 41 BP; 8 A; 13 C; 12 G; 8 T; 0 U; 0 Other;

S Matches Query Match Best Local Similarity 643 CCCAGGCTGGAGTGCAGTGGCGCAATCTTGGCTCACTGCAA 683 Conservative 3.5%; 0 Score 34.6; Pred. No. 3 Mismatches 3.7e+02 DВ 1; Length 4; Indels 41; 0 Gaps 0

ADL64139 standard; DNA; 41 BP

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41

CCCAGGCTGGAGTGCAGTGGTGCGATCTCAGCTCACTGCAA 1

single nucleotide polymorphism (SNP)

20-MAY-2004

(first entry)

RESULT 207
ADL64139
ID ADL641
XX ADL641
XX DT 20-MAN
XX Y
DT 20-MAN
XX S8; h
KW C1 S; k
KW C1 S s
KW tissum
KW tissum
KW solubKW solubKW solubKW solubKW cardic
KW myocaa
KW myocaa
KW myocaa
KW myocaa
KW myocaa
KW cancei
KW asthmm
KW cancei
KW Asthme
KW Asthme ss; human; single nucleotide polymorphism; SNP; C1 S subcomponent protein; C1S; alanyl aminopeptidase protein; ANPEP; meprin A beta protein; aminopeptidase P-like protein; XPN-PEPL; tissue kallikrein protein; KIKI, aminopeptidase P protein; MEPIB; soluble guanylate cyclase 1 alpha-2 subunit protein; GUCY1A2; haplotype; angioedema; angioedema-like disorder; paternity testing; cardiovascular disease; angina pectoris; hypertension; heart failure; myocardial infarction; aneurysm; stroke; embolism; thrombosis; coronary artery disease; arteriosclerosis; hypersensitivity; haemodialysis; sepsis; inflammatory disease; inflammatory arthritis; haemodialysis; sepsis; inflammatory disease; inflammatory disease; inflammatory disease. chronic

Homo sapiens

WPI; 2004-180052/17.

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                                                                                                                                                        cc human gene encoding a protein, such as the C1, S subcomponent protein CC (C1S), the alanyl aminopeptidase protein (ANPP), the meprin A, beta C protein (MEP1B), the aminopeptidase P-like protein (XPN-PEPL), the tissue CK stalikrein protein (KUKI), the membrane bound aminopeptidase P protein (CC (SUCYIA2). The nucleic acid comprises at least one polymorphic position, cincluding the alleles, reference alleles and alternate alleles of the CC single nucleotide polymorphisms, listed in the specification. The CC single nucleotide polymorphisms, listed in the specification within the genomic sequence of the gene. The polymorphic position residing in a coding position results in a missense or silent mutation of the translated CC product of the gene. The polymorphic position residing in a non-coding position resides within the untranslated region or an intronic region of the gene. Constructing haplotypes using the nucleic acids above further CC of a disease phenotype, and correlating the presence of the disease phenotype with the haplotype. The disease phenotype is angioedema or an angioedema-like disorder. The nucleic acids, primers and probes are CC useful in phenotype correlations, paternity testing, medicine and genetic snalysis. The nucleic acids and polypeptides can be used in diagnosing, coronary acternosclerosis, hypersensitivity reactions during haemodialysis, constructive pulmonary diseases, inflammatory arthritis, asthma, chronic constructive pulmonary diseases, inflammatory allergies, or cancer. The nucleotion for the presence of the constructive pulmonary diseases, inflammatory allergies, or cancer. The nucleotion mucleotion during haemodialysis, the pulmonary diseases, inflammatory arthritis, asthma, chronic constructive pulmonary diseases, inflammatory allergies, or cancer. The nucleotide polymorphism, constructive pulmonary diseases, inflammatory arthritis, asthma, chronic constructive pulmonary diseases, inflammatory arthritis, asthma, chronic constructive pulmonary diseases, inflammatory arthritis, a
                                                                              Matches
                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New nucleic acid comprising a single nucleotide polymorphism at a specific location, useful in paternity testing, genetic analysis or diagnosing, preventing or treating cardiovascular diseases e.g.
                                                                                                                                                      Sequence 41
                                                                                                                                                                                                               present sequence represents a human single nucleotide polymorphism
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 3; SEQ ID NO 62; 376pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      angioedema or angina pectoris.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       invention relates to an isolated nucleic acid (I) derived from
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RAMANATHAN C
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                        AGCTGGGACCAAAGACATGCACCACTACACCTGGCTAATTT 597
AGCTGGGATTACAGACATGCACCACCACCTGGCTAATTT 41
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. Z, Zerba
                                                                                                                                                    BP; 12
                                                                            Conservative
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                                                                                            3.5%;
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Pred. No. 3.7e+02;
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                                                                          Mismatches
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                                                                          4.
                                                                                                            Length
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                                                                          Gaps
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ss; human; single nucleotide polymorphism; SNP; cl S subcomponent protein; ANPEP; alanyl aminopeptidase protein; ANPEP; meprin A beta protein; minopeptidase P Pilke protein; MEPIB; tissue kallikrein protein; KLKI; aminopeptidase P protein; MEPIB; soluble guanylate cyclase 1 alpha-2 subunit protein; GUCYIA2; haplotype; angioedema; angioedema-like disorder; paternity testing; cardiovascular diseases; angina pectoris; hypertension; heart failure; myocardial infarction; aneurysm; stroke; embolism; thrombosis;
                                                                                                                                                                                                                                                                                  Edmonds M,
Tsuchihashi
                                                                                                                                                                                                                                                                                                                                                        (EDMO/)
(HUIL/)
(PERR/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cancer; ANPEP.
                                                                                                                                                                                                                                                                                                                               (RAMA/)
(SWAN/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       coronary artery disease; arteriosclerosis; hypersensitivity; haemodialysis; sepsis; inflammatory disease; inflammatory ar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human single nucleotide polymorphism
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                                                                                                                                                                                                                                                                                                              (ZERB/)
                                                                                                                                                                                                                                                                                                                                                 (POWE/)
                                                                                                                                                                                                                                                                                                              ZERBA
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RAMANATHAN C S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               chronic obstructive pulmonary disease;
                                                                                                                                                                                                                                                                                                                               SWANSON B.
                                                                                                                                                                                                                                                                                                                                                                            EDMONDS
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Z,
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ba K;
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                                                                                                                                                                                                                                                                                            Ramanathan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cough reflex; allergy;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        arthritis;
                                                                                                                                                                                                                                                                                             Swanson
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NEW NUCLEIC acid comprising a single nucleotide polymorphism specific location, useful in paternity testing, genetic analy diagnosing, preventing or treating cardiovascular diseases e. angioedema or angina pectoris. analysis õ

WPI; 2004-180052/17.

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Claim 3; SEQ ID NO 207; 376pp; English.

CC human gene encoding a protein, such as the C1. S subcomponent protein CC (C1S), the alanyl aminopeptidase protein (ANPEP), the meprin A, beta CC protein (MEP1B), the aminopeptidase Prike protein (XPN-PEPL), the tissue CC (Allikrein protein (KIXI), the mebrane bound aminopeptidase P protein (XPNPEP2), or the soluble guanylate cyclase 1, alpha-2 subunit protein CC (GUCY1A2). The nucleic acid comprises at least one polymorphic position, CC including the alleles, reference alleles and alternate alleles of the Single nucleotide polymorphisms, listed in the specification. The CC single nucleotide polymorphisms, listed in the specification. The CC golymorphic position resides in a (non)coding position within the genomic Sequence of the gene. The polymorphic position residing in a coding CC position resides within the untranslated comprises using the haplotypes using the nucleic acids above further CC comprises using the haplotypes to identify an individual for the presence of a disease phenotype, and correlating the presence of the disease correlations, paternity testing, medicine and genetic analysis. The nucleic acids and polypeptides can be used in diagnosing, The invention relates ç an isolated nucleic acid (I) derived from s subcomponent protein

RESULT 208 ADL64284/c

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RESULT 209
AAT97406/c
Query Match
Best Local S
Matches 37
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                                            A method has been developed for detecting the presence of a target site (TS), of at least one nucleotide (nt) in a nucleic acid (NA). The method comprises: (a) forming an oligonucleotide (ON), consisting of two fluorophores (F1, F2) each covalently linked to separate nt, bound to TS; and (b) detecting fluorescence energy transfer (FET) between F1 and F2 when ON is released from TS. The present sequence represents a synthetic polynucleotide used in an example of the present invention. The method is used to diagnose hereditary and other diseases; to determine infectious agents; in tissue typing for histocompatibility; in forensic identification and paternity testing, and in monitoring the genetic make up of plants and animals. Specifically it is used to detect single nt polymorphisms. The method provides inexpensive, simple, accurate and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 41 BP; 7 A; 14 C;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     present sequence represents a human single nucleotide polymorphism
                                                                                                                                                                                                                                                                                                                          Detecting target site in nucleic acid by forming a fluorophore-labelled oligonucleotide at the site - and detecting fluorescent energy following denaturation, used e.g. to detect inherited diseases, in tissue typing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Detection; target site; nucleic acid; fluorophore; labelled; fluorescent; inherited disease; tissue typing; PCR; ss.
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                                                                                                                                                                                                                                                                          Example 2; Page 27; 68pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                   Kwok P,
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                                   acid analyses
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No. 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 1; Length 41;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
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Query Match

Sequence

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33.6; Ή, 0

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1; Length Other;

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Sequence 40

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RESULT 210
ADK41334/c
ID ADK413
The invention relates to a novel method of estimating disease risk or cc prognosis of an individual by sequence polymorphism analysis, especially conjumorphisms in the human chromosome 19q. The invention further relates to estimating a treatment response of an individual suffering from cc cancer to a disease treatment; a primer or probe for use in the method of cestimating the disease risk or prognosis of an individual or for cestimating treatment; an antibody directed to an epitope of a RAI gene conjumors, and a kit for use in the method of estimating the disease risk or prognosis of an individual suffering the disease risk or prognosis of an individual or for estimating the disease risk or prognosis of an individual or for estimating a treatment, comprising at cleast one primer or probe and optionally amplifying means for nucleic carid amplification. The novel method is useful for estimating the disease risk or prognosis of an individual or for estimating a treatment response of an individual suffering from cancer to a disease treatment response of an individual suffering from cancer to a disease treatment response of an individual suffering from cancer to a disease treatment. This
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                27-JUN-2002; 2002DK-00001005.
07-OCT-2002; 2002DK-00001500.
25-FEB-2003; 2003DK-00000289
29-APR-2003; 2003DK-00000639
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADK41334
                                                                                                                                                                                                                                                                                                                                                             Estimating the disease polymorphism analysis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       single nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                          Nexo BA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  27-JUN-2003; 2003WO-DK000448
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                08-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             variation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       sequence polymorphism analysis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human chromosome 19 single nucleotide polymorphism detecting
                                                                                                                                                                                                                                                                                                                             Claim 18; SEQ ID NO 92; 145pp; English
                                                                                                                                                                                                                                                                                                                                                          polymorphism
                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2004-142878/14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO2004003229-A2
                                               polynucleotide sequence represents a probe used nucleotide polymorphisms in the DNA of human chr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (ARBE-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (UYAA-)
                                            nucleotide polymorphisms in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              675
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             36;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        UNIV AARHUS.
ARBEJDSMILJO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TCACTGCAACCTCTGCCTCCCGGGTTCAAGTTATTCTCCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TCACTGCAAGCTCTGCCTCCTGGGTTCAAGCAATTCTCCT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                         Vogel U,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers replace(20,G)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /*tag= a
/standard_name= "Single nucleotide polymorphism"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     polymorphism; SNP; probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA:
                                                                                                                                                                                                                                                                                                                                                                                                                                          Rockenbauer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            90.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INST NAT INST
                                                                                                                                                                                                                                                                                                                                                                            risk or prognosis of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pred. No. 4.1
), Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           human;
                                                                                                                                                                                                                                                                                                                                                                                                                                          'n
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OCCUPA
                                                                                                                                                                                                                                                                                                                                                                                                                                            Bukowy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           chromosome 19q; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                            X
                                                                                                                                                                                                                                                                                                                                                                               an
                                               chromosome
                                                                                                                                                                                                                                                                                                                                                                               individual by sequence
                                               for detecting single romosome 19 of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 714
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          probe #22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RAI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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RESULT 212
ABL49776
ID ABL497
XX ABL497
AC ABL497
XZ ABL497
XX DT 29-MAY
XX TX
DT Human
XX THUMAN
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Best Local S
                                                                                                                                                                                                                                                                                                                                                                                          Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                              Human tyrosinase 10.34 probe 2
                                                                                             29-MAY-2002
                                                                                                                                        ABL49776;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 7; Page 20; 36pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA mismatch repair protein 11 and encoded polynucleotide, diagnosis and treatment of malignant tumor, hemopathy, HIV immunological diseases and various inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mao Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (UYFU-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               23-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       18-DEC-2000; 2000WO-CN000627.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   05-JUL-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200147988-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; DNA repair mismatch protein 11; cancer; haemopathy; HIV infection; immunological disease; inflammation; gene therapy; probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human DNA mismatch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAH49728
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAH49728 standard;
                                                                                                                                                                                  ABL49776 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention provides the protein and coding sequences of human DNA mismatch repair protein 11. The sequences are useful in the treatment of cancer, haemopathy, HIV infection, immunological diseases and of cancer, the present sequence is a probe for the coding sequence of the coding seque
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sapiens.
                                                                                                                                                                                                                                                                                                                                        839
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2001-425639/45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     469
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                                                                                                                                                                                                                                                                                                41
  tyrosinase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        UNIV FUDAN.
SHANGHAI BI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                        TCTGCCTGCCTCGGCCTCCCAAAGTGCTGGGATTACAGGC 878
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CCCAGGCTGGAGTGCAGTGCGATCTCAGCTCACTGCA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CCCAGGATGAAGTGCAGTGGTGATCACAGCTCACTGCA 508
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP; 9 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                           (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               99CN-00125733.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BIO DOOR GENE TECHNOLOGY LTD
  enzyme; human immunodeficiency virus infection;
                                                                                                                                                                                  DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  repair protein 11 coding sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA;
                                                                                                                                                                                                                                                                                                                                                                                                          3.4%;
                                                                                         entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            11 C; 15 G; 6
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                                                                                                                                                                                  BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0,
                                                                                                                                                                                                                                                                                                                                                                                     <u>,,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 33.6; DB Pred. No. 4.1e+
                                                                                                                                                                                                                                                                                                                                                                                                          Score 33.6; DB 1;
Pred. No. 4.2e+02;
                                              SEQ
                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              T; 0 U;
                                                 ID NO:9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                               Length 41;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 40;
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                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                infection,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       applicable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         #2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
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                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
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RESULT 213
ABL49775
ID ABL497
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention describes human tyrosinase 10.34 (I). The present invention also described a method for preparing (I) using DNA recombination techniques. (I) and the polymuleotide encoding it can be used in the treatment of diseases such as cancer and human immunodeficiency virus (HIV) infection. The present sequence represents probe for human tyrosinase 10.34, which is used in an example from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mao
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HIV infection;
                                                 Mao
                                                                                          31-MAY-2000;
                                                                                                                                                       CN1325972-A.
                                                                                                                                                                                              Human; tyrosinase; enzyme; human HIV infection; cancer; probe; ss.
                                                                                                                                                                                                                            Human tyrosinase 10.34 probe 1 SEQ ID NO:8.
                                                                                                                                                                                                                                                  29-MAY-2002
                                                                                                                                                                                                                                                                      ABL49775;
                                                                                                                                                                                                                                                                                          ABL49775
                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 41 BP; 5 A; 12 C; 13 G; 11 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 6; Page 19 (Disclosure); 32pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New polypeptide-tyrosinase 10.34 for treating human immunodeficiency virus infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 31-MAY-2000; 2000CN-00116261
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       31-MAY-2000; 2000CN-00116261
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CN1325972-A.
                             WPI; 2002-196693/26.
                                                                                                             31-MAY-2000; 2000CN-00116261
                                                                                                                                   12-DEC-2001.
                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12-DEC-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (BODE-)
                                                 ۲,
                                                                                                                                                                                                                                                                                                                                                                  187
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                                                 Xie
                                                                     BODE
                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
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                                                                                                                                                                                                                                                                                                                                                           TGGAGTTTCTCCATGTTGGTCAGGCTGGTCTCGAACTCCC 226
                                                                                                                                                                                                                                                                                                                                              TGGGGTTTCACCATGTTGGCCGGGCTGGTCTCGAACTCCC
                                                                                                                                                                                                                                                                                           standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                      GENE
                                                                                           2000CN-00116261
                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cancer; probe; ss.
                                                                      DEV
                                                                                                                                                                                                                                                                                                                                                                                                3.4%;
                                                                     8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CO LTD SHANGHAI
                                                                                                                                                                                                                                                                                           41
                                                                      GLT
                                                                                                                                                                                                                                                                                           ВP
                                                                                                                                                                                                                                                                                                                                                                                      <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                Score 33.6; DB 1
Pred. No. 4.2e+02
                                                                      SHANGHAI.
                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                        immunodeficiency virus infection;
                                                                                                                                                                                                                                                                                                                                                                                                         DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  diseases such
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New polypeptide-tyrosinase 10.34 for to human immunodeficiency virus infection.

for treating

diseases

such

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cancer and

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RESULT 214
AREZOLG 7/C
ID ABZ206
XX ABZ206
XX ABZ206
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XX O3-MAR
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Matches 36
                                                                                          Matches
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; G protein subunit 9.02; cardiac asthma; colic; psychic morphinic analgesic acute poisc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABZ20667 standard;
                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                              cancers,
                                                                                                                                                                                                                                                                                                                                                                        Example
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (SHAN-) SHANGHAI BIOWINDOW GENE DEV INC.
                                                                                                                                                                                                                           he present invention provides the protein and coding sequences of human protein subunit 9.02. The sequences can be used in the treatment of ancers, coughs, cardiac asthma, diarrhoea, constipation, colic, psychic isease and morphinic analgesic acute poisoning. The present sequence is probe used to isolate the coding sequence of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    187
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                                                                                                                                                                                                                                                                                                                                                                     6; Page 22 (Disclosure); 34pp; Chinese.
                                                                                                                  Similarity
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  TCCACCCACCTCGGCCTCCCGAAGTGCTGGGATTACAGGC
                                         TCCACCTGCCTCAGCCTCCCAAAGTGCTGGGATTACAGGC 408
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                                                                                                                                                                                    BP;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       entry)
                                                                                                               3.4%;
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                                                                                                                                                                                    10 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      41
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                                                                                                                                                                                    16 G; 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 33.6; DB 1;
Pred. No. 4.2e+02;
                                                                                          Score 33.6; DB 1;
Pred. No. 4.2e+02;
0; Mismatches 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cancer; constipation; diarrhoea; cough;
disease; probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               11
                                                                                                                                                                                    T; 0 U; 0 Other;
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                                                                                                                                        DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Other;
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                                                                                                                                     Length 41;
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SXXXXXXXXXXXXX

Human oncogene

protein 11-66 nucleotide probe

19-SEP-2002

(first entry)

Homo sapiens inflammation;

Human; ss; gene therapy; haemopathy; development

probe

gene therapy; oncogene protein development disturbance; HIV;

11.66; malignant tumou
immunological disease;

tumour;

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RRESULT 215
ARBA94091/C
ID ARBA940
XX ABA940
XX ABA940
XX ABA940
DT 08-MAY
XX Human
XX Human
XX Human
XX Human
XX Homo s
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CC (I) he
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CC HIV)
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CC (II) he
CC (III)
                    RESULT 216
AAL43826
                                                                                                                                                                                               Query Match
Best Local S
Matches 36
                                                                                                                                                                                                                                                                                                                              The present invention describes human tumour suppressor factor 11.77 (1). (1) has cytostatic, haemostatic, virucide, immunomodulatory and antiinflammatory activities. The polynucleotide (II) encoding (I) can be used in gene therapy. (I) and (II) can be used in the diagnosis and used the second of the diagnosis and the diagnosis and the second of malignant tumour, haemopathy, human immunodeficiency virus (HIV) infection, immunological diseases and various inflammations. The present sequence represents a probe for human tumour suppressor factor 11.77, which is used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human tumor suppressor factor 11.77 and encoding polynucleotide, diagnosis and treatment of malignant tumors, hemopathy, human immunodeficiency virus infection, immunological diseases and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      immunomodulatory; antiinflammatory; gene therapy; malignant tumour;
haemopathy; human immunodeficiency virus infection; HIV infection;
immunological disease; inflammation; probe; ss.
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                                                                                                                                                                                                                                                                                    Sequence 41 BP; 9 A; 11 C; 14 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 6; Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2002-075588/10.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20-DEC-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; tumour suppressor factor 11.77;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  inflammation.
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                                                                                                                                                        839
                                                                                                                                                                                                    36;
                                                                                                            41
                                                                                                                                                                                                                        Similarity
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standard;
                                                                                                       TCCGCCCGTCTTGGCCTCCCAAGTGCTGGGATTACAGGC
                                                                                                                                     TCTGCCTGCCTCGGCCTCCCAAAGTGCTGGGATTACAGGC
                                                                                                                                                                                                      Conservative
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    DNA; 41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      33pp; Chinese.
                                                                                                                                                                                                                        3.4%;
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Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DEV
                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                             33.6; DB 1
No. 4.2e+02
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                                                                                                                                                                                                                                                  DB 1;
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Best Local S
Matches 36
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mao
                                                                                                                                                                                                                                         Human oncogene
                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 6; Page 21 (Disclosure); 33pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel polypeptide-oncoprotein 11.66\, and polynucleotide polypeptide.
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                           polypeptide
                                                                                                            07-JUL-2000; 2000CN-00119427.
                                                                                                                              07-JUL-2000; 2000CN-00119427
                                                                                                                                                30-JAN-2002
                                                                                                                                                                  CN1333235-A
                                                                                                                                                                                                              Human; ss; gene therapy; oncogene protein 11.66; malignant haemopathy; development disturbance; HIV; immunological dis
                                                                                                                                                                                                                                                           19-SEP-2002
                                     Novel polypeptide-oncoprotein 11.66 and
                                                                                                                                                                                                       inflammation;
                                                                                                                                                                                                                                                                                              AAL43827 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (SHAN-)
                                                                                           (SHAN-)
                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
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                                                                                                                                                                                    sapiens.
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                                                                                           SHANGHAI
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                                                                                                                                                                                                                                                                                                                                                             TGGAGTTTCTCCATGTTGGTCAGGCTGGTCTCGAACTCCC
                                                                                                                                                                                                                                                                                                                                                                                                                  BP; 6 A;
                                                                                                                                                                                                                                                                                                                                                                               Conservative
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                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                      probe
                                                                                                                                                                                                                                         protein 11-66 nucleotide probe
                                                                                           BIODOOR GENE
                                                                                                                                                                                                                                                                                              DNA; 41
                                                                                                                                                                                                                                                                                                                                                                                                                   13
                                                                                                                                                                                                                                                                                                                                                                                       3.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                  C; 12
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Pred. No. 4.2e+02;
0; Mismatches 4
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                                     polynucleotide for encoding
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Example 6; Page

21

(Disclosure);

33pp;

Chinese.

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RESULT 218
ABZ44551/c
ID ABZ445
XX ABZ44551/c
AC ABZ445
XX ABZ445
XX ABZ445
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Best Local (
Sequences ABZ43217-ABZ50887 represent polymorphic sites within genes encoding enzymes associated with drug metabolism. The invention relates to methods and compositions for identifying individuals who have at least one polymorphism in such drug metabolising enzyme-encoding genes. The polymorphisms may be identified in a nucleic acid sample using probes or primers specific for a sequence selected from ABZ43217-ABZ50887 using a variety of detection assays, including hybridisation assays, nucleic acid arrays and PCR-based methods. The invention also encompasses methods of evaluating and screening drugs using genetic polymorphism data, particularly that relating to single nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention comprises the amino acid and coding sequence of the human oncogene protein 11.66. The oncogene protein 11.66 DNA and protein sequences are useful for treating malignant tumour, haemopathy, development disturbance, HIV infection, immunological disease and various inflammations. The present DNA sequence represents a probe that is specific for the gene sequence of the human oncogene protein 11.66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; drug metabolising enzyme; gene; drug metabolism; chromo polymorphic site; drug evaluation; drug screening; genotyping; genetic profiling; therapeutic customisation; adverse reaction clinical trial; drug approval; single nucleotide polymorphism;
                                                                                                                                                                                                                                                                                                                                                                                                   Identifying individuals having a polymorphism, useful for determining effectiveness or side effect of a drug or treatment protocol, comprise detecting at least one polymorphism in the drug metabolizing enzyme
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       27-DEC-2000;
02-MAY-2001;
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                                                                                                                                                                                                                                                                                                                      23; Page 86; 2785pp; English.
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2001JP-00135256.
2001JP-00256862.
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Pred. No. 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      reaction;
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polymorphisms (SNPs), may be used in studying the relationship between DNA sequence variations and human diseases, conditions, and responses

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RESULT 219
ABZ50761/c
ID ABZ5077
XX ABZ5077
XX ABZ507
XX Human
XX Human
XX Human
XX Human
XX Homo s
XX Clinic
XX Cli
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                                                                                                                                                                                                                      27-DEC-2000; 2000JP-00399443
02-MAY-2001; 2001JP-00135256
27-AUG-2001; 2001JP-00256862
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                                    WPI; 2002-583571/62
                                                                                                 Nakamura Y,
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                                                                                                    Sekine
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/*tag= a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /standard_name= "Single nucleotide polymorphism (SNP)"
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Pred. No. 4
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Identifying individuals having a polymorphism, useful for determining the effectiveness or side effect of a drug or treatment protocol, comprises detecting at least one polymorphism in the drug metabolizing enzyme
                                                                                                                                                                                                  detecting at least nucleic acid.
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23; Page 221; 2785pp; English

CC sequences ABZ43217-ABZ50887 represent polymorphic sites within genes CC encoding enzymes associated with drug metabolism. The invention relates CC one polymorphism in such drug metabolising enzyme-encoding genes. The CC one polymorphisms may be identified in a nucleic acid sample using probes or CC primers specific for a sequence selected from ABZ43217-ABZ50887 using a CC variety of detection assays, including hybridisation assays, mucleic acid carrays and PCR-based methods. The invention also encompasses methods of CC evaluating and screening drugs using genetic polymorphism data. particularly that relating to single nucleotide coloryphism data, particularly that relating to single nucleotide polymorphisms (SNPs), may be used in studying the relationship between cc polymorphisms (SNPs), may be used in studying the relationship between cc polymorphisms (SNPs), may be used in studying the relationship between cc polymorphisms (SNPs), may be used in studying the relationship between cc that cause or exacerbate certain diseases, SNPs are particularly useful critical cause or exacerbate certain diseases, SNPs are particularly useful critical cause or exacerbate certain diseases, SNPs are particularly useful critical trials based upon the genetic profile of individual patients. CC drug therapias based upon the genetic profile of individual patients of this would not only take the genetic profile of individual patients. CC methods of the invention are also useful in the drug discovery and capable of the invention are also useful in the drug discovery and capable of responding to a particular functions, thereby increasing safety. CC capable of responding to a particular discovery capable of responding to a particular discovery capable of the invention may therefore lead to a an increase in the range of the invention may therefore lead to a an increase in the range of the invention may therefore the definition and the number of different contributions. The methods, data and compositions of the length of this patient medications a patient needs to take before finding an effective

Sequence 41 BP; 10 A; 8 C; 16 G; 7 T; 0 U; 0 Other;

Query Match Best Local S Matches 36 36; Similarity Conservative 3.4%; 0, Score 33.6; Pred. No. 4 Mismatches DB 1; Length 0; Gaps 0

669 CTTGGCTCACTGCAACCTCTGCCTCCCGGGTTCAAGTTAT 708

밁 á RESULT 220 ABV77329 standard; DNA; 41 40 CTTGGCTCACTGCAACCTCCGCCTCCTGGGTTCAAGCAAT 1 ВP

Human protein 10.01 related probe 2.

07-FEB-2003 ABV77329;

(first entry)

Human; 10.01; aminolyase active site; arrhythmia; diabetes; probe; 83

CN1342770-A

12-SEP-2000; 2000CN-00125186

12-SEP-2000; 2000CN-00125186

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RESULT 221
ACC00157
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Best Local
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         present
protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         polynucleotide, and a method for preparing the polypeptide by DNA recombination. The application of the polypeptide is in treating arrhythmia and diabetes. Also disclosed are the antagonist against this polypeptide and its therapeutic action, and the application of the polypeptide. The current sequence represents a human protein 10.01 related noche sequence.
                                                                                                               A polypeptide-guanosine triphosphatase activator polynucleotide for coding this polypeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        ACC00157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to a human protein designated 10.01, containing Phe-His aminolyase active site. Also disclosed are the encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example
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V
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mao
                               The present invention discloses a polypeptide-guanosine triphosphatase activator protein-10.01. The invention also discloses the method for curing several diseases, such as squamobasal cell carcinoma of skin, osteosarcoma, leukemia and teratoma by using said polypeptide. The
                                                                                                                                                                                                                                                                                                 CN1380320-A.
                                                                                                                                                                                                                                                                                                                        Unidentified
                                                                                                                                                                                                                                                                                                                                                carcinoma
                                                                                                                                                                                                                                                                                                                                                          Guanosine
                                                                                                                                                                                                                                                                                                                                                                                 Probe #2 for guanosine triphosphate activator 10.01.
                                                                                                                                                                                                                                                                                                                                                                                                          14-JUL-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                ACC00157;
                                                                                          Example 7; Page 23; 33pp; Chinese.
                                                                                                                                                                                                                           10-APR-2001; 2001CN-00105912.
                                                                                                                                                                                                                                                10-APR-2001; 2001CN-00105912
                                                                                                                                                                                                                                                                          20-NOV-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   related probe sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (BODE-)
                                                                                                                                                                                                   (SHAN-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    human protein 10.01 oding polynucleotide,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ۲
                                                                                                                                                    2003-222552/22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         655
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               36;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  _
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Xie
                                                                                                                                                                            Xie
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BODE GENE
           sequence represents a 10.01
                                                                                                                                                                                                   SHANGHAI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              41
                                                                                                                                                                                                                                                                                                                                                                                                                                                       standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TGCAGTGGCGCAATCTTGGCTCACTGCAACCTCTGCCTCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TGCAGTGGCGCAATCTTGGTCCACTGCAACCCCCCCCCTCC
                                                                                                                                                                                                                                                                                                                                              얁
                                                                                                                                                                                                                                                                                                                                                          triphosphatase activator 10.01; squamobasal cell;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP; 6 A; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                              skin;
                                                                                                                                                                                                                                                                                                                                                                                                          (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22
                                                                                                                                                                                                   BIOWINDOW GENE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DEV
                                                                                                                                                                                                                                                                                                                                                osteosarcoma; leukemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (disclosure); 33pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                         entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ç
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  containing Phe-His aminolyase active site and useful for treating arrhythmia and diabetes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 33.6; DB 1;
Pred. No. 4.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SHANGHAI.
                                                                                                                                                                                                   DEV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                              teratoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 41;
                                                                                                                            protein -10.01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         694
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                   polypeptide. The triphosphate activator
                                                                                                                                                                                                                                                                                                                                              probe; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                              and
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RESULT 223
ADL64136/c
ID ADL641
XX
AC ADL641
XX
DT 20-MAY
XX

ADL64136; ADL64136

standard;

DNA;

41 ВP

20-MAY-2004

(first entry)

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RESULT 222
ABZ57114
ID ABZ571
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                    S
                                                          Best
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local
                                              Matches
                                                                    Query Match
                                                                                                                         The invention relates to human KIAA0608 protein 10.12 (ABP58674) and nucleic acids encoding it (ABZ57108). The protein has a molecular weight of 10 kD. The invention also relates to a method for the recombinant production of the protein, an antagonist of the protein, and the use of the protein, gene and antagonist in therapeutic applications. KIAA0608 protein 10.12 can be used in the treatment of a variety of diseases such as peptic ulcers and diabetes. Sequences ABZ57113-ABZ57114 represent human KIAA0608 protein 10.12 probes used in an exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 41
                                                                                                                                                                                                                                                                                                         Mao
                                                                                            Sequence 41 BP; 6 A; 11 C;
                                                                                                                                                                                                                                    Example
                                                                                                                                                                                                                                                                                  WPI; 2003-000145/01.
                                                                                                                                                                                                                                                                                                                                                                                                                       CN1355220-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; KIAA06
peptic ulcer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human KIAA0608 protein 10.12 probe, SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABZ57114;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABZ57114
                                                                                                                   invention
                                                                                                                                                                                                                                                          Polypeptide-human KIAA0608 protein 10.12 and polynucleotide encoding
                                                                                                                                                                                                                                                                                                                                                      24-NOV-2000; 2000CN-00127565
                                                                                                                                                                                                                                                                                                                                                                            24-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                   26-JUN-2002
                                                                                                                                                                                                                                                                                                                               (UYFU-) UNIV FUDAN
                                                         Local
                       648 GCTGGAGTGCAGTGGCGCAATCTTGGCTCACTGCAACCTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              655
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    36;
                                              36;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 KIAA0608 protein 10.12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       L
N
                                                                                                                                                                                                                                                                                                         Xie
                                                          Similarity
                                                                                                                                                                                                                                   6; Page 22 (Disclosure); 35pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TGCAGTGGCGCAATCTTGGCTCACTGCAACCTCTGCCTCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TGCAGTGGCACAGTCTCGGCGCACTGCAACCTCTGCCTCC
 GCTGGAGTGCAGTGACGCAATCTTGGCTCGCTGCAGGCTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP; 6 A;
                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                             2000CN-00127565
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      diabetes; probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16 C; 11 G; 8 T; 0 U; 0 Other;
                                                         90.0%;
                                                                     3.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              41
                                                                                            14 G; 10 T; 0 U; 0 Other;
                                              0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 33.6;
Pred. No. 4
                                                          Pred. No. 4.2e+02
                                                                     Score 33.6; DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         .88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  recombinant production;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NO: 9
                                               4.
                                                                    Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               694
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                         687
                                               Indels
  41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  gene therapy;
                                                                      41;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            41;
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                                              0
                                              Gaps
                                                                                                                                                                                                                                                           it.
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CC human gene encoding a protein (AURED), the meprin A, beta CC (CIS), the alanyl aminopeptidase P-like protein (XPN-BPI), the tissue CC (Rallikrein protein), the membrane bound aminopeptidase P protein (XPN-BPI), the tissue CC (Rallikrein protein), the tissue CC (Rallikrein protein), the membrane bound aminopeptidase P protein CC (GUCYIA2). The nucleic acid comprises at least one polymorphic position, cincluding the alleles, reference alleles and alternate alleles of the cCC single nucleotide polymorphisms, listed in the specification. The CC sequence of the gene. The polymorphic position residing in a coding position resides in a missense or silent mutation of the translated CC position resides within the untranslated region or an intronic region of the gene. Constructing haplotypes using the nucleic acids above further CC comprises using the haplotypes to identify an individual for the presence CC of a disease phenotype, and correlating the presence of the disease phenotype is angioedema or an angioedema -like disorder. The nucleic acids, primers and probes are CC useful in phenotype correlations, paternity testing, medicine and genetic CC analysis. The nucleic acids and polypeptides can be used in diagnosing, preventing or treating cardiovascular disease, e.g. angioedema, angina pectoris, hypertension, heart failure, myocardial infarction, aneurysm, stroke, embolism, thrombosis, coronary artery disease or cancer. The obstructive pulmonary disease, cough reflex, allergies, or cancer. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              meprin A beta protein; aminopeptidase P-like protein; XPM-PEPL; tissue kallikrein protein; KIKI; aminopeptidase p protein; MEPIB; soluble guanylate cyclase 1 alpha-2 subunit protein; GUCYIA2; haplotype; angioedema; angioedema-like disorder; paternity testing; cardiovascular diseases; angina pectoris; hypertension; heart failure; myocardial infarction; aneurysm; stroke; embolism; thrombosis; coronary artery disease; arteriosclerosis; hypersensitivity; haemodialysis; sepsis; inflammatory disease; inflammatory arthritis; asthma; chronic obstructive pulmonary disease; cough reflex; allergy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human single nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         specific location, useful in patern diagnosing, preventing or treating angioedema or angina pectoris.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New nucleic acid comprising a single nucleotide polymorphism at a specific location, useful in paternity testing, genetic analysis of diagnosing, preventing or treating cardiovascular diseases e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     03-JUN-2003; 2003US-00453827
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US2004033582-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cancer; ANPEP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             03-JUN-2002; 2002US-0384980P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19-FEB-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to an isolated nucleic acid (I) derived from a human gene encoding a protein, such as the Cl, S subcomponent protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tsuchihashi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (RAMA/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EDMO/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ຜ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZERBA K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RAMANATHAN C S
SWANSON B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   POWELL J R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PERRONE M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EDMONDS M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TSUCHIHASHI Z.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OI DES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     single nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hui L,
Z, Ze
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NO 59; 376pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Zerba
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    uncleotide polymorphism; SNP; protein; ClS; alanyl aminono-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Perrone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             polymorphism
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Σ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Powell JR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (SNP)
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P-like protein; XPN-PEPL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ramanathan
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RESULT 224
ABZ45510
IID ABZ45510
IID ABZ45510
ABZ45510
AC ABZ455
XX ABZ455
XX ABZ455
XX Human
XX Human
XX Human
XX Human
XX Human
XX Clinic
XX Clinic
XX WO2000
YX Claim
XX Ident
YX Claim
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CC PI mucle
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CC One polym
CC One polym
CC Dolym
CD D
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one polymorphism in such drug metabolising enzyme-encoding genes. The polymorphism may be identified in a nucleic acid sample using probes or primers specific for a sequence selected from ABZASONET using a variety of detection assays, including hybridisation assays, nucleic acid arrays and PCR-based methods. The invention also encompasses methods of evaluating and screening drugs using genetic polymorphism data, particularly that relating to single nucleotide polymorphisms (SNPs), may be used in studying the relationship between DNA sequence variations and human diseases, conditions, and responses to drugs. SNPs are also useful as polymorphism markers for discovering genes that cause or exacerbate certain diseases. SNPs are particularly useful
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02-MAY-2001; 2001JP-00135256
27-AUG-2001; 2001JP-00256862
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                                                                                                                                                                                                                                                                                                          Sequences ABZ43217-ABZ50887 represent polymorphic sites within genes encoding enzymes associated with drug metabolism. The invention relates to methods and compositions for identifying individuals who have at least
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human ATP-binding cassette ABCA7 gene polymorphic site,
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Pred. No. 4.2e
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RESULT 225
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Best Local Similarity
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02-MAY-2001;
27-AUG-2001;
Identifying individuals having a polymorphism, useful for determining the effectiveness or side effect of a drug or treatment protocol, comprises detecting at least one polymorphism in the drug metabolizing enzyme
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Pred. No. 4.3e+02;
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CC encoding enzymes associated with drug metabolism. The invention relates come polymorphism in such drug metabolising enzyme-encoding genes. The compositions for identifying individuals who have at least cone polymorphisms may be identified in a nucleic acid sample using probes or compilers specific for a sequence selected from ABZ4217-ABZ50887 using a covariety of detection assays, including hybridisation assays, nucleic acid arrays and PCR-based methods. The invention also encompasses methods of covariety of detection assays, including hybridisation assays, nucleic acid carrays and PCR-based methods. The invention also encompasses methods of covaluating and screening drugs using genetic polymorphism data. Genetic colymorphisms (SNPs), may be used in studying the relationship between compasses to drugs. SNPs are also useful as polymorphism markers for discovering genes to that cause or exacerbate certain diseases. SNPs are particularly useful in the above respects as they are stable in populations, and responses to that cause or exacerbate certain diseases. SNPs are particularly useful cin penes encoding drug metabolising enzymes allows the customisation of the genes of cover mutation rates than other genome variations of the invention are also useful an populations, occur in genes encoding drug metabolising enzymes allows the customisation of creduce the likelihood of adverse reactions, thereby increasing safety.

CC methods of the invention are also useful in the drug discovery and spiral processes. For example, individuals could be selected for approval processes. For example, individuals could be selected for capropriate partient populations. The methods, data and compositions of the invention may therefore lead to an increase in the range of the invention spiral trials are on methods, data and compositions of the invention are also methods. The number of adverse drug to be approved, the length of time patients are on medication and the number of different medications a patient people.
                                                                                                      Sequence '41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequences ABZ43217-ABZ50887
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 23; Page 129; 2785pp; English.
                                                                                                           B₽;
                                                                                                           8 A;
                                                                                                                 14 C; 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               represent polymorphic sites within genes with drug metabolism. The invention rela
                                                                                                           G; 10 T; 0 U; 0 Other;
DB 1; Length 41;
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밁 á Matches Query Match POCST 667 4 l Similarity 35; Conserv ATCTTGGCTCACTGCAACCTCTGCCTCCCGGGTTCAAG 704 ATCTTGGCTCACTGCAACCTCCGCCTCCTGGATTCAAG Conservative 3.4%; <u>,</u> Score 33.2; Pred. No. 4. Mismatches 4.3e+02; 41 Indels 0 Gaps

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19-JUL-2001;
19-JUL-2001;
16-NOV-2001;
                                                                                             Cytostatic; Antibacterial; Immunosuppressive; neural thread protein; NTP; tumour; ds.
                                                                                                                   NTP peptide
                                     19-JUL-2002;
                                                                                Unidentified.
                                                                                                                                  28-AUG-2003
                                                                                                                                                              ACC84461 standard;
                                                                  WO2003008443-A2
                                                                                                                   encoding sequence
        2001US-0306150P.
2001US-0306161P.
2001US-0331477P.
                                                                                                                                  (first entry)
                                     2002WO-CA001105
                                                                                                                                                               DNA;
                                                                                                                                                               ü
                                                                                                                     #8
                                                                                                      Antiinflammatory;
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nucleic

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RESULT 227
AAI99796
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Best Local S
Matches 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to a neural thread protein (NTP) peptide referred to as cell death peptide. Thought to be cytostatic, antibacterial, immunosuppressive and antiinflammatory. It is useful for treating a condition in a patient requiring removal or destruction of cells, for treating a condition such as benign or malignant tumor, inflammatory disease, autoimmune disease and infectious disease. The peptide useful for treatment is derived from the amino acid sequence for a pancreatic thread protein. The peptide is conjugated, linked or bound to a molecule chosen from antibody or its fragment, antibody-like binding molecule, where the molecule has a higher affinity for binding to a tumor or other target than binding to other cells. Treatment using NTP peptides can remove benign tumors with less risk and fewer of the undesirable side effects of surgery. The present sequence is an NTP encoding sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 17; 77pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel neural thread protein peptide, referred as cell death peptide, useful for treating prostatic hyperplasia, psoriasis, eczema, dermatosis, atherosclerosis, cosmetic modification to skin, throat, mouth, muscle.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               P-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Averback PA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             XOMAN (-OWAN)
              Human eucaryotic acetyl transferase 10 and encoded polynucleotide, in diagnosis and treatment of malignant tumors, hemopathy, human immunodeficiency virus infection, immunological diseases and
                                                                                                                                                                                                                                                                                                                                                                                                         Human eukaryotic acetyl transferase 10 probe SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                          24-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAI99796 standard;
                                                                                                                                                                                                                                                                 WO200175026-A2
                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                           Human; eukaryotic acetyl transferase 10; cytostatic; virucidal;
                                                                                                                                                                     22-MAR-2000; 2000CN-00115031
                                                                                                                                                                                                   19-MAR-2001; 2001WO-CN000378
                                                                                                                                                                                                                                                                                                                                              human
                                                                                                                                                                                                                                                                                                                                            immunomodulatory; antiinflammatory; haemostatic; malignant
numan immunodeficiency virus; HIV; infection; immunological
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   378
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2003-247999/24.
                                                                                                                                                                                                                                                                                                                            therapy; probe; ss
                                                                             2002-025848/03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABR63256.
                                                                                                          Xie
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                         BIOWINDOW GENE DEV INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CTCAGCCTCCCAAAGTGCTGGGATTACAGGCGT 410
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CTCAGCCTCCCAAAGTGCTGGGATTACAGGCGT 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP; 7 A; 10 C; 9 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ₽₽
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                                                                                                                                         SHANGHAI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB: 1;
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                                                                                                                                                                                                                                                                                                                                                tumour;
l disease;
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RESULT 228
ABL52956/c
ID ABL52956
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Best Local S
Matches 36
                                                         Query Match
Best Local :
                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to human eukaryotic acetyl transferase 10 with cytostatic, virucidal, immunomodulatory, antiinflammatory and haemostati activity. The protein and encoding polynucleotide are used in diagnosis and treatment of malignant tumour, haemopathy, human immunodeficiency virus (HIV) infection, immunological diseases and various inflammations. The polynucleotide is useful in gene therapy. The present sequence is that of a probe, useful to the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 6; Page 15; 33pp; Chinese
                                                                                                                               The present invention relates to serine proteinase 10 (AAM48453). Serine proteinase 10 and its coding sequence can be used for treating diseases such as cancer and HIV infection. The present sequence is a probe, which
                                                                                                                                                                                                                                                                                                                                                                                                                                      Serine proteinase 10; enzyme; cancer; HIV infection; anti-HIV; cytostatic; probe; ss.
                                                                                                                                                                                                         Polypeptide-serine proteinase 10 and polynucleotide encoding
                                                                                                                                                                                                                                                                                                                                                                                                                Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Serine proteinase 10 probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    24-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABL52956;
                                                                                             Sequence 41 BP; 9 A; 15 C; 12 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                   Example 6;
                                                                                                                                                                                                                                    WPI; 2002-196715/26.
                                                                                                                                                                                                                                                                                                             31-MAY-2000; 2000CN-00116278
                                                                                                                                                                                                                                                                                                                                     31-MAY-2000;
                                                                                                                                                                                                                                                                                                                                                               12-DEC-2001.
                                                                                                                                                                                                                                                                                                                                                                                       CN1325996-A.
                                                                                                                       was used
                                                                                                                                                                                                                                                                                    (BODE-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1051 TGCCACCACACCCCGCTAATTTTTGTATTTTCATTAGAGGC 1091
                  637 CTGTCACCCAGGCTGGAGTGCAGTGGCGCAATCTTGGCTCA 677
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   36;
41
                                                36;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                      BODE GENE
                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TGCCACCACGCCGGCTAATTTTTGTATTTTTAGTAGAAGC
                                                                                                                      in an example
                                                                                                                                                                                   Page 19 (Disclosure); 32pp; Chinese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                       2000CN-00116278
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                      DEV CO LTD SHANGHAI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA;
                                                         3.3%;
                                                                                                                         from the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        41
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Pred. No.
                                                           Score
Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                 Mismatches
                                                           33;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4.4e+02;
                                                           4.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1;
                                                 5; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 41;
                                                                       Length 41;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      41
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                                                 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            haemostatic
                                                 Gaps
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RESULT 229 ABZ44124/c

0

ABZ44124 standard; DNA; 41

ВP

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Sequences ABZ43217-ABZ50887 represent polymorphic sites within genes concoding enzymes associated with drug metabolism. The invention relates cone polymorphism in such drug metabolising enzyme-encoding genes. The polymorphism is such drug metabolising enzyme-encoding genes. The cone polymorphism is such drug metabolising enzyme-encoding genes. The primers specific for a sequence selected from ABZ43217-ABZ50887 using a curaity of detection assays, including hybridisation assays, nucleic acid carrays and PCR-based methods. The invention also encompasses methods of CC evaluating and screening drugs using genetic polymorphism data. Genetic polymorphism data, particularly that relating to single nucleotide conformations and numan diseases. SNPs are lationship between CC polymorphism data, particularly that relating to single nucleotide conformations and numan diseases. SNPs are particularly genes to CC drugs. SNPs are also useful as polymorphism markers for discovering genes that cause or exacerbate certain diseases. SNPs are particularly useful circular cause or exacerbate certain diseases. SNPs are particularly useful circular states than other genome variations cour frequently, and have lower mutation rates than other genome variations cour genes encoding drug metabolising enzymes allows the customisation of drug therapies based upon the genetic profile of individual patients. CC frist would not only take the guessork out of selecting the drug with the greatest therapeutic effect for a particular patient, but would also course the likelihood of adverse reactions, thereby increasing safety. CC Methods of the invention are also useful in the drug discovery and secure and secure and secure and form of the guessor of the invention may therefore lead to an increase in the range of the invention of decreases in the methods, data and compositions of the invention may therefore lead to a mincrease in the range of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     27-DEC-2000;
02-MAY-2001;
27-AUG-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; drug metabolising enzyme; gene; drug metabolism; chromosc
polymorphic site; drug evaluation; drug screning; genotyping;
genetic profiling; therapeutic customisation; adverse reaction;
clinical trial; drug approval; single nucleotide polymorphism;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Identifying individuals having a polymorphism, useful for determining effectiveness or side effect of a drug or treatment protocol, comprise detecting at least one polymorphism in the drug metabolizing enzyme
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nakamura
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          27-DEC-2001;
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drug targets and
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2001JP-00135256.
2001JP-00256862.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers replace(21,T)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         polymorphic
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re lead to a an increase in decreases in the number of
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RESULT 23
RABZ45508
ID ABZ45508
ID ABZ4
XX ABZ4
AC ABZ4
AC ABZ4
AC ABZ4
AC ABZ4
XX BOLL
AC ABZ4
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02-MAY-2001;
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l trial; drug approval; single nucleotide
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Pred. No. 4.
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Sequences ABZ43217-ABZ50887 represent polymorphic sites within genes encoding enzymes associated with drug metabolism. The invention relates to methods and compositions for identifying individuals who have at least one polymorphism in such drug metabolising enzyme-encoding genes. The polymorphisms may be identified in a nucleic acid sample using probes or primers specific for a sequence selected from ABZ43217-ABZ50887 using a variety of detection assays, including hybridisation assays, nucleic acid arrays and PCR-based methods. The invention also encompasses methods of evaluating and screening drugs using genetic polymorphism data, particularly that relating to single nucleotide polymorphisms (SNPs), may be used in studying the relationship between DNA sequence variations and human diseases, conditions, and responses to detecting at least Identifying individuals having a polymorphism, useful for determining the effectiveness or side effect of a drug or treatment protocol, comprises 23; Page 102; one polymorphism in 2785pp; English or treatment protocol, the drug metabolizing enzyme ecid

are also useful as polymorphism markers

and responses to discovering gener

genes

Nakamura Y,

Sekine

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Saito

(RIKE) RIKEN KK.

Claim 23; Page 192; 2785pp; English.

nucleic acid

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RESULT 231
ABZ49572/C
ID ABZ495
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ID ABZ495
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ABZ49
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02-MAY-2001;
27-AUG-2001;
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Identifying individuals having a polymorphism, useful for determining effectiveness or side effect of a drug or treatment protocol, comprise detecting at least one polymorphism in the drug metabolizing enzyme
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Pred. No.
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ches 5;
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CC sequences ABZ4317-ABZ50887 represent polymorphic sites within genes CC encoding enzymes associated with drug metabolism. The invention relates CC one polymorphism in such drug metabolising enzyme-encoding genes. The cone polymorphisms may be identified in a nucleic acid sample using probes or CC primers specific for a sequence selected from ABZ4327-ABZ50887 using a cC variety of detection assays, including hybridisation assays, nucleic acid carrays and PCR-based methods. The invention also encompasses methods of CC evaluating and screening drugs using genetic polymorphism data. Genetic CC polymorphism data, particularly that relating to single nucleotide colony sequence variations and human diseases, conditions, and responses to CC MANA sequence variations and human diseases. SNPs are particularly useful colony in the above respects as they are stable in populations, occur crequently, and have lower mutation rates than other genome variations cc in genes encoding drug metabolising enzymes allows the customisation of creduce the likelihood of adverse reactions, thereby increasing safety. This would not only take the genesure reactions, thereby increasing safety. Methods of the invention are also useful in the drug discovery and capable of responding to a particular partient, but would also approval processes. For example, individuals could be selected for capable of responding to a particular drug or drug class, and previously cfailed drug candidates could be revived if they were matched with more capacitions, failed drug trials, the time taken for a drug to be approved, the length of time patients are on medication and the number of different could be selected for a possible drug targets and decreases in the number of adverse of the therapy of the patients are on medication and the number of different the lands of the selections of the length of time patients are on medication and the number of different the patient is not take before finding an effective therapy the could be selected for an interest of diff
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  RESULT 232
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Best Local S
Matches 36
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variation
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Sequences ABZ43217-ABZ50887 represent polymorphic sites within genes encoding enzymes associated with drug metabolism. The invention rela
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Sequences ABZ43217-ABZ50887 represent polymorphic sites within genes cancoding enzymes associated with drug metabolism. The invention relates to methods and compositions for identifying individuals who have at least one polymorphism in such drug metabolising enzyme-encoding genes. The polymorphisms may be identified in a nucleic acid sample using probes or primers specific for a sequence selected from ABZ43217-ABZ50887 using a covariety of detection assays, including hybridisation assays, nucleic acid arrays and PCR-based methods. The invention also encompasses methods of carrays and PCR-based methods. The invention also encompasses methods of collusting and screening drugs using genetic polymorphism data. particularly that relating to single nucleotide polymorphisms (SNPs), may be used in studying the relationship between CC polymorphisms (SNPs), may be used in studying the relationship between CC polymorphisms useful as polymorphism markers for discovering genes to drugs. SNPs are also useful as polymorphism markers for discovering genes chat cause or exacerbate certain diseases. SNPs are particularly useful cc in the above respects as they are stable in populations, occur
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02-MAY-2001; 2001JP-00135256
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Identifying individuals having a polymorphism, useful for determining effectiveness or side effect of a drug or treatment protocol, comprise detecting at least one polymorphism in the drug metabolizing enzyme nucleic acid.

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2002-583571/62

Claim 23;

76; 2785pp; English.

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                                                                                                                                                                                                                                                                            Human; drug metabolising enzyme; gene; drug metabolism; chromosome polymorphic site; drug evaluation; drug screening; genotyping; genetic profiling; therapeutic customisation; adverge reaction; clinical trial; drug approval; single nucleotide polymorphism; SNP;
WO200252044-A2
                                                                                           variation
                                                                                                                                                               variation
                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                Human ATP-binding cassette ABCA7
                                                                                                                                                                                                                                                                                                                                                                                                                                                26-JUN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABZ46914 standard; DNA; 41 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    reactions, failed drug trials, the time taken for a drug to be approved, the length of time patients are on medication and the number of different medications a patient needs to take before finding an effective therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      361 TCAAGCAGTCCACCTGCCTCAGCCTCCCAAAGTGCTGGGAT 401
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                                                                               /standard name= "Single nucleotide polymorphism replace(25,T)
/*t-n
                                                                                                                                                          Location/Qualifiers replace(21,A)
                                      standard_name= "Single nucleotide polymorphism (SNP)"
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Pred. No. 4.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                             gene polymorphic site,
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                                                                                                                 (SNP) "
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CC variety of detection assays, including hybridisation assays nucleic acid contrarys and PCR-based methods. The invention also encompasses methods of CC evaluating and screening drugs using genetic polymorphism data. Genetic polymorphism data, particularly that relating to single nucleotide cCC polymorphism (SNPs), may be used in studying the relationship between CC drugs. SNPs are also useful as polymorphism markers for discovering genes that cause or exacerbate certain diseases, conditions, and responses to frequently, and have lower mutation rates than other genome variations cCC in genes encoding drug metabolising enzymes allows the customisation of CC drug therapies based upon the genetic porofile of individual patients. CCC This would not only take the guesswork out of selecting the drug with the greatest therapeutic effect for a particular patient, but would also creduce the likelihood of adverse reactions, thereby increasing safety. CC capable of responding to a particular drug or drug discovery and CC capable of responding to a particular drug or drug class, and previously failed drug targets and decreases in the methods, data and compositions of cups that they are particular genetic profiles indicate that they are appropriate patient populations. The methods, data and compositions of the invention may therefore lead to a an increase in the range of compositions failed drug trials, the time taken for a drug to be approved, the length of time patients are as methods and the number of different reads to take before finding an effective therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             27-DEC-2000; : 02-MAY-2001; : 27-AUG-2001; :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            sequences ABZ43217-ABZ50887 represent polymorphic sites within genes encoding enzymes associated with drug metabolism. The invention relates to methods and compositions for identifying individuals who have at least one polymorphism in such drug metabolising enzyme-encoding genes. The polymorphisms may be identified in a nucleic acid sample using probes or primers specific for a sequence selected from ABZ43217-ABZ50887 using a variety of detection assays, including hybridisation assays.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Identifying individuals having a polymorphism, useful for determining the effectiveness or side effect of a drug or treatment protocol, comprises detecting at least one polymorphism in the drug metabolizing enzyme
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                                         medications a patient needs to take before finding an effective
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2001JP-00135256.
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                      RESULT 237
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Best Local S
Matches 36
ABZ47296 standard; DNA;
                                                                                                                                                         Sequence 41 BP; 6
                                                                              639 GTCACCCAGGCTGGAGTGCAGTGGCGCAATCTTGGCTCACT 679
                                                                                                           36;
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                                                                                                                       Similarity
                                                                                                                                                         A,
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                                                                                                          Score 33; DB 1; Length 41
Pred. No. 4.4e+02;
0; Mismatches 5; Indels
                                                                                                                                  Length 41;
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therapy

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CC encoding enzymes associated with drug metabolism. The invention relates cc to methods and compositions for identifying individuals who have at least cone polymorphism in such drug metabolising enzyme-encoding genes. The polymorphism in such drug metabolising enzyme-encoding genes. The cc primers specific for a sequence selected from ABZ43217-ABZ50887 using a cc variety of detection assays, including hybridiaation assays, mucleic acid carrays and pCR-based methods. The invention also encompasses methods of cc evaluating and screening drugs using genetic polymorphism data, particularly that relating to single nucleotide cc polymorphism (aNPs), may be used in studying the relationship between cc polymorphisms (SNPs), may be used in studying the relationship between cc polymorphisms (SNPs), may be used in studying the relationship between cc palymorphisms (SNPs), may be used in studying the relationship between cc respects as they are stable in populations, and responses to compasses methods of the above respects as they are stable in populations, occur frequently, and have lower mutation rates than other genome variations such mass repeating sequences. The detection and analysis of polymorphisms cc in genes encoding drug metabolising enzymes allows the customisation of creduce the likelihood of adverse reactions, their patient, but would also creduce the likelihood of adverse reactions, thereby increasing safety. Cc methods of the invention are also useful in the drug discovery and apreviously creasing to a particular drug or drug class, and previously falled drug candidates could be revived if they were matched with more capable drug candidates could be revived if they were matched with more capactons, failed drug trials only trials on medication and the number of different medications a patient needs to take before finding an effective therapy of the length of time patients are on medication and the number of different medications a patient needs to take before finding an effective therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Identifying individuals having a polymorphism, useful for determining the effectiveness or side effect of a drug or treatment protocol, comprises detecting at least one polymorphism in the drug metabolizing enzyme
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nakamura Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         27-DEC-2000; 2000JP-00399443
02-MAY-2001; 2001JP-00135256
27-AUG-2001; 2001JP-00256862
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (RIKE ) RIKEN KK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Page 138;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sekine
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Iida A,
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RESULT 238
ABA94080
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Best Local Similarity
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                                                     Matches
                                                                Query Match
Best Local (
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                                                                                                                                              has cytostatic, haemostatic, virucide, immunomodulatory and antiinflammatory activities. The polynucleotide (II) encoding (I) can be used in gene therapy. (I) and (II) can be used in the diagnosis and treatment of malignant tumour, haemopathy, human immunodeficiency virus (HIV) infection, immunological diseases and various inflammations. The present sequence represents a probe which is used in an example from the
                                                                                                                                                                                                                                                                                       Multi-copper oxidase 12 and encoding polynucleotide, used in diagnos and treatment of malignant tumors, hemopathy, human immunodeficiency virus infection, immunological diseases and inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; multi-copper oxidase 12; enzyme; cytostatic; haemost virucide; immunomodulatory; antiinflammatory; gene therapy; malignant tumour; haemopathy; human immunodeficiency virus HIV infection; immunological disease; inflammation; probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human multi-copper oxidase 12 probe 1 SEQ ID NO:8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    08-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABA94080;
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                                                                                                                                                                                                                                                             Example 7; Page 14; 31pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                WPI; 2002-075593/10.
                                                                                                                                                                                                                                                                                                                                                                          мао У,
                                                                                                                                                                                                                                                                                                                                                                                                                              19-MAY-2000; 2000CN-00115756
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20-DEC-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                       Sequence 41 BP; 6 A; 6 C; 14 G; 15 T; 0 U; 0 Other;
                                                                                                                                                                                                                                The present
                                                                                                                                                                                                                                                                                                                                                                                                    (BIOW-) BIOWINDOW GENE DEV INC SHANGHAI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         369 TCCACCTGCCTCAGCCTCCCAAAGTGCTGGGATTACAGGCG 409
                           172
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                                                                                                                                    invention
                                                                   Similarity
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                         TTTTTTAGTAGAGATGGAGTTTCTCCATGTTGGTCAGGCT 212
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                   invention describes human multi-copper oxidase 12
                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA; 41
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                                                                  3.3%;
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Pred. No.
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                                                      Mismatches
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                                                        Indels
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                                                                                                                                                                                                                                                                                                                        in diagnosis
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RESULT 239 AAS15590

AAS15590

AAS15590 standard; DNA; 41 BP

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  repair protein 10. The polypeptide can be used for screening mimics, agonists, antagonists or inhibitors, or in peptide fingerprinting identification. The polymucleotide can be used as primers for nucleic acid amplification reactions, as probes for hybridisation reactions, or in producing gene chips or microarrays. Drug compositions, which contain the polypeptide, polymucleotide, mimics, agonists, antagonists, inhibitors and their preparations, can be used treatment and diagnosis of diseases relating to the polypeptide. In particular, the polypeptide and encoded polynucleotide are applicable. In diagnosis and treatment of malignancy, haemopathy, human immunodeficiency virus (HIV) infection, immunological diseases and various inflammations. The present sequence
ss; human; single nucleotide polymorphism; SNP; C1 S subcomponent protein; C1S; alamyl aminopeptidase protein; ANPEP; meprin A beta protein; aminopeptidase P like protein; XPN-PEPL; tissue kallikrein protein; KLKI; aminopeptidase P protein; MEPIB; soluble guanylate cyclase 1 alpha-2 subunit protein; GUCY1A2; haploty
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; DNA mismatch repair protein 10; cytostatic; virucidal; immunomodulatory; antiinflammatory; haemostatic; anti-HIV; inflammation; human immunodeficiency virus; malignancy; haemopathy; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New human DNA mismatch repair protein 10 for diagnosing and treating malignancy, hemopathy, human immunodeficiency virus infection, immunological diseases and inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14-FEB-2002
                                                                                                      Human single
                                                                                                                                        20-MAY-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 41 BP; 8 A; 10 C; 11 G; 12 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    immunological diseases and various inflammations. The present sequence represents probe #1 used in Northern blot analysis of human DNA mismatch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       repair protein
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                                                                                                                                                                                                                                                                                                                                                                                   Similarity
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                                                                                                                                                                                                        standard; DNA;
                                                                                                                                                                                                                                                                                                                                 TGTTGGTCAGGCTGGTCTCGAACTCCCGACCTCAGATGATC 240
                                                                                                                                                                                                                                                                                                TGTTGGTCAGACTGGTCTTGAACTCCCAACGTCAGGTGATC 41
                                                                                                                                                                                                                                                                                                                                                                   Conservative
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                                                                                                    nucleotide polymorphism (SNP)
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                                                                                                                                        (first
                                                                                                                                        entry)
                                                                                                                                                                                                                                                                                                                                                                               3.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                an isolated polypeptide of human DNA mismatch
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                                                                                                                                                                                                                                                                                                                                                          Pred. No. 4.4c
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                 Score 33;
Pred. No.
 subunit protein; GUCY1A2; haplotype;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     probe
                                                                                                                                                                                                                                                                                                                                                                                 4.4e+02
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                                                                                                      #208.
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                                                                                                                                                                                                                                                                                                                                                                                                   Length 41;
                                                                                                                                                                                                                                                                                                                                                                   Indels
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comparison the alanyl aminopeptidase protein (ANPER), the malanyl aminopeptidase protein (ANPER), the membrane bound aminopeptidase Portein (XPN-PEPL), the tissue (CRIS), the alanyl aminopeptidase Portein (ANPER), the meprin A, beta (CRIS), the alanyl aminopeptidase Portein (XPN-PEPL), the tissue (CRIS), the meprin A, beta (CRIS), the meprin A (CRIS), the membrane bound aminopeptidase protein (CRIS), the alleles, reference alleles and alternate alleles of the crisical polymorphic position and alternate alleles of the crisical polymorphic position residing the polymorphic position residing the polymorphic position residing in a non-coding (CRIS), the gene. The polymorphic position residing in a non-coding (CRIS), the gene. Cristophenotype of polymorphic position residing in a non-coding (CRIS) and the presence of the gene. Cristophenotype, and correlating the nucleic acids above further (CRIS) and the haplotypes to identify an individual for the presence of a disease phenotype, and correlating the presence of the disease phenotype is anyloedema or an angioedema-like disorder. The nucleic acids, primers and probes are useful in phenotype correlations, paternity testing, medicine and genetic analysis. The nucleic acids and polypeptides can be used in diagnosing, percorris, hypertension, heart failure, myocardial infarction, aneurysm, cristophenotype in the membrane or anteriosclerosis, hypersensitivity reactions during haemodialysis, espesia, inflammatory diseases, cough reflex, allergies, or cancer. The presente a human single nucleotide polymorphism (SNP) of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New nucleic acid comprising a single nucleotide polymorphism at specific location, useful in paternity testing, genetic analysis diagnosing, preventing or treating cardiovascular diseases e.g. angioedema or angina pectoris.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cardiovascular diseases; angina pectoris; hypertension; heart myocardial infarction; aneurysm; stroke; embolism; thrombosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Edmonds M,
Tsuchihashi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (EDMO/)
(HUIL/)
(PERR/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        coronary artery disease; arteriosclerosis; hypersensitivity; haemodialysis; sepsis; inflammatory disease; inflammatory ar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to an isolated nucleic acid (I) derived from human gene encoding a protein, such as the Cl, S subcomponent prote
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 3; SEQ ID NO 208; 376pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2004-180052/17.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     asthma; chronic obstructive pulmonary disease; cough reflex; allergy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           angioedema; angioedema-like disorder; paternity testing;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (rama/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (POWE/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZERBA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SWANSON B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RAMANATHAN C S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TSUCHIHASHI Z.
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Z,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Perrone
ba K;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 S,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                heart failure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       arthritis;
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Query Match Best Local Similarity

3.3%;

Score 33; Pred. No.

DB 1;

Length 41;

Sequence

41

BP; 5 A;

14 C; 11 G; 11 T; 0 U; 0 Other;

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RESULT 24
RESULT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ss; human; single nucleotide polymorphism; SNP;
Cl S subcomponent protein; ClS; alanyl aminopeptidase protein; ANPEP;
meprin A beta protein; aminopeptidase P-like protein; XPN-PEPL;
tissue kallikrein protein; KLKL; aminopeptidase P protein; MEPIB;
soluble guanylate cyclase 1 alpha-2 subunit protein; GUCY1A2; haplotype;
angioedema; angioedema-like disorder; paternity testing;
cardiovascular diseases; angina pectoris; hypertension; heart failure;
myocardial infarction; aneurysm; stroke; embolism; thrombosis;
The invention relates to an isolated nucleic acid (I) derived from a human gene encoding a protein, such as the Cl, S subcomponent protein (CIS), the alanyl aminopeptidase protein (ANPEP), the meprin A, beta protein (MEPIB), the aminopeptidase P-like protein (XPN-PEPL), the tissue kallikrein protein (KUKI), the membrane bound aminopeptidase P protein (XPNPEP)), or the soluble guanylate cyclase 1, alpha-2 subunit protein (GUCYLA2). The nucleic acid comprises at least one polymorphic position, including the alleles, reference alleles and alternate alleles of the single nucleotide polymorphisms, listed in the specification. The polymorphic position resides in a (non)coding position within the genomic sequence of the gene. The polymorphic position residing in a coding position results in a missense or silent mutation of the translated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human single nucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New nucleic acid comprising a single nucleotide polymorphism at a specific location, useful in paternity testing, genetic analysis or diagnosing, preventing or treating cardiovascular diseases e.g.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Edmonds M,
Tsuchihashi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cancer; ANPEP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 angioedema or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2004-180052/17
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SWANSON B.
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                                                                                                                                                                                                                                                                                                                                                                                         SEQ ID NO
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Z,
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Zerba
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Powell JR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ramanathan CS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               5
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Identifying individuals having a polymorphism, useful for determining effectiveness or side effect of a drug or treatment protocol, comprise detecting at least one polymorphism in the drug metabolizing enzyme nucleic acid.

WPI; 2002-583571/62

Nakamura Y, (RIKE) RIKEN

Sekine

Iida

P

Saito

즛

Claim 23; Page 168; 2785pp; English

27-DEC-2000; 2000JP-00399443: 02-MAY-2001; 2001JP-00135256. 27-AUG-2001; 2001JP-00256862.

27-DEC-2001; 2001WO-JP011592

04-JUL-2002. WO200252044-A2 variation

Location/Qualifiers replace(20. .21,GGT)

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88888888888888888888888888888888888
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ABZ48532
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 242
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                position resides within the unitranslated region or an intronic region of the gene. Constructing haplotypes using the nucleic acids above further comprises using the haplotypes to identify an individual for the presence of a disease phenotype, and correlating the presence of the disease compensation of the disease phenotype with the haplotype. The disease phenotype is angioedema or an angioedema-like disorder. The nucleic acids, primers and probes are considered in phenotype correlations, paternity testing, medicine and genetic analysis. The nucleic acids and polypeptides can be used in diagnosing, correcting or treating cardiovascular diseases, e.g. angioedema, angina conjunctions, heart failure, myocardial infarction, aneurysm, stroke, embolism, thrombosis, coronary artery disease or arteriosclerosis, hypersensitivity reactions during haemodialysis, cobstructive pulmonary disease, inflammatory arthritis, asthma, chronic obstructive pulmonary disease, cough reflex, allergies, or cancer. The present sequence represents a human single nucleotide polymorphism (SNP)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      product of the gene. The polymorphic position residing in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human, drug metabolising enzyme; gene; drug metabolism; polymorphic drug evaluation; drug screening; genotyping; genetic profiling; therapeutic customisation; adverse reaction; clinical trial;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human oligopeptide transporter PEPT1 gene polymorphic site, #5315
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 41 BP; 11 A; 13 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                grug
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          26-JUN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABZ48532;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABZ48532 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                approval;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    557 AGCTGGGACCAAAGACATGCACCACTACACCTGGCTAATTT 597
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                36;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AGCTGGGATTACAGACATGCCCCACCACACCTGGCTAATTT 41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          entry!
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           8 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 33;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 41;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             a non-coding
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Compared the likelihood of adverse reactions, thereby increasing safety. This would not only take the guesswork out of selecting the drug with the greatest therapeutic effect for a particular patient, but would also concesses. For example, individuals could be selected for clinical trials only if their genetic profiles indicate that they are appropriate patient populations. The methods, data and compositions of the invention may therefore lead to a mincrease in the range of compositions of the invention synthericals they were matched with more appropriate patient populations. The methods, data and compositions of the length of time patients are on medication and the number of different medications a patient needs to take before finding an effective therapy.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 243
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local S
Matches 35
                                                                                                                                                                                                                                                                   Yeast Artificial Chromosome; YAC; polymerase chain reaction; PC sequence tagged site; genetic disorder; diagnosis; abnormality; Prader-Willi; Angelman; Beckwith-Wiedermann; syndrome; ss.
                                                                                                                                                                                                                                                                                                                                                                                     25-MAR-2003
28-OCT-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                             AAQ45257;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAQ45257 standard; DNA; 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    one polymorphism in such drug metabolising enzyme-encoding genes. The polymorphisms may be identified in a nucleic acid sample using probes or primers specific for a sequence selected from ABZ43217-ABZ50887 using a variety of detection assays, including hybridisation assays, nucleic aci
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequences ABZ43217-ABZ50887 represent polymorphic sites within genes encoding enzymes associated with drug metabolism. The invention relato methods and compositions for identifying individuals who have at
   Airhart SD,
                                                                             11-SEP-1992;
                                                                                                                                                        31-MAR-1994
                                                                                                                                                                                             WO9406936-A1
                                                                                                                                                                                                                                                                                                                                                Alu primer PDJ34 to amplify Yeast Artificial Chromosome DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      medications a patient needs to take before finding an effective therapy
                                       (BAYU )
                                                                                                                  10-SEP-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
nes 35; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1076
                                       BAYLOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TATTTTCATTAGAGGCGGGGTTTCACCATATTTGTCAGGC 1115
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TATTTTTAGTAGAGACGGGGTTTCACCATATTGGCCAGGC 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP; 9 A; 7 C; 11 G; 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                   (revised)
(first en
 Mutirangura A, Ledbetter DH;
                                       COLLEGE MEDICINE
                                                                             92US-00943639.
                                                                                                                  93WO-US008501.
                                                                                                                                                                                                                                                                                                                                                                                     entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ₽₽
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RESULT 244
ABA93847/c
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
          The present invention describes human GASC1 (gene amplified in squamous cell carcinoma 1). GASC1 has been located to the p23-24 region of human chromosome 9. GASC1 can be used in the diagnosis and investigation of diseases with which cell differentiation and proliferation are associated, such as cancer. It can also be used in gene therapy of these diseases; and screening substances for their ability to modify the expression of GASC1 and for use as drugs. The present sequence represents a PCR primer for human GASC1, which is used in an example from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   particular, probes to diagnose Prader-Willi/Angelman Syndrome were identified. The majority of PWS/Angelman patients are deleted for a common set of markers including ML34, IR4-3R, TD189-1 and TD3-21. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           amplify DNA from yeast artificial chromosomes as part of a cloning procedure to isolate probes for specific chromosomal abnormalities.
                                                                                                                                                                                      Gene GASC1 amplified in squamous cell carcinoma and its expression product for diagnosis investigation and treatment of disorders inve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cell proliferation; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; GASC1; gene amplified in squamous cell carcinoma 1; cancer; chromosome 9; chromosome 9p23-24; cell differentiation; gene thera:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human GASC1 PCR primer SEQ ID NO:5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           02-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABA93847
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Diagnosis of genetic disorders associated with chromosomal abnormalities and uniparental disomy, e.g. Prader-Willi:Angelman syndrome - using in
                                                                                                                                             Example 1; Page
                                                                                                                                                                                                                                 WPI; 2002-090209/12
                                                                                                                                                                                                                                                              Inazawa
                                                                                                                                                                                                                                                                                                                      12-JUN-2000; 2000JP-00174946.
                                                                                                                                                                                                                                                                                                                                                   12-JUN-2001; 2001WO-JP004959
                                                                                                                                                                                                                                                                                                                                                                                20-DEC-2001.
                                                                                                                                                                                                                                                                                                                                                                                                             WO200196566-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens.
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                                                                                                                                                                                                                                                                                          (SAKA)
                                                                                                                                                                         proliferation
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27; Conser
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                                                                                                                                                                                                                                                                                           OTSUKA PHARM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5; Page 32; 91pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           primers PDJ34 and 2484 (AAQ45257 and AAQ45258) were used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CCCAGGCTGGAGTGCAGTGGCGCAATCTTGGCTCA 677
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     standard;
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                                                                                                                                                                                                                                                              Imoto I;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first
                                                                                                                                          77; 82pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA;
                                                                                                                                                                           such as cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          entry
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Pred. No. 4.5e
8; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0 U; 8 Other,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    gene therapy;
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             represents
                                                                                                                                                                                         involving
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present invention

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RESULT 246
ADE14248/c
ID ADE142
XX
AC ADE142
XX
AC ADE142
XX
29-JAN
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AAQ27392/c
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Best Local (
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                                                                                                                                                   Query Match
Best Local S
                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAQ27392 standard; DNA;
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                                                                                                                                                                                                     Primer PDJ34 is one of several primers which are preferred for use in amplifying inter-Alu regions of DNA. The amplified fragments are then subjected to 2-D electrophoresis on the basis of length and differences in base sequence. The resulting separation pattern is transferred to a filter for screening with a probe. The method can be used to detect genetic variation. See AAQ27389-Q27404 and AAQ33141-Q33144. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                           WPI;
                                                                                                                                                                                                                                                                                                              Detection of genetic variation by 2-D electrophoresis of fragments hybridisation with labelled probes, carried out on fragments consi of inter-repeat sequences generated by PCR.
                                                                                                                                                                                                                                                                                                                                                                                Uitterlinden AG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      06-AUG-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO9213101-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25-MAR-2003
27-JAN-1993
                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                            25-JAN-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                24-JAN-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Polymerase chain reaction; PCR; repetitive element; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Inter-Alu
  29-JAN-2004
                                             ADE14248
                                                                                                                                                                                                                                                                                            Claim 6; Page 6; 31pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
                                                                                                                                                                                                                                                                                                                                                             1992-284683/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              643
                                                                                                                    643
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                                                                                                 35
                                                                                                                                        l Similarity
32; Conserv
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5
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                                                                                                                  CCCAGGCTGGAGTGCAGTGGCGCAATCTTGGCTCA 677
                                             standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         specific primer
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                                                                                                                                                                                   BP; 6 A;
                                                                                                                                         Conservative
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  (first entry)
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                                                                                                                                                  3.2%;
                                             DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                entry)
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                                                                                                                                                                                   12 C; 11 G; 5 T; 0.U; 1 Other;
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                                             32
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                                             ВÞ
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B; Mismatches
                                                                                                                                                   Score 31.4; DB 1;
Pred. No. 4.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score
                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         31.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               .5e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               677
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
                                                                                                                                                              Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 35;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                          Indels
                                                                                                                                                                35;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     <u>,</u>
                                                                                                                                         0
                                                                                                                                                                                                                                                                                                                             consisting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
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SNP; glaucoma; progressive ocular hypertensive disorder;
glaucoma related disorder; motif; repeat element; regulatory region.
                                                                                                                                                                                                                                                                                                                                   Human; optineurin;
                                                                                                                                                                                                                                                                                                                                                                        Optineurin promoter motif,
                               Raymond V,
                                                                                                                                  06-MAR-2002; 2002US-00091281
                                                                                                                                                                     06-MAR-2002; 2002US-00091281
                                                                                                                                                                                                                                     US2003190617-A1.
                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                 (SIEE/)
                                                                 (MORI/) MORISSETTE
2003-864168/80
                                                                                 SI E.
RAYMOND V.
                               Morissette J,
                                                                                                                                                                                                                                                                                                                                          ds; ophthalmological; single
                                                                                                                                                                                                                                                                                                                                                                           repeat element
                                   Si
                                 Ħ
                                                                                                                                                                                                                                                                                                                                                                             or regulatory
                                                                                                                                                                                                                                                                                                                                           nucleotide polymorphism;
                                                                                                                                                                                                                                                                                                                                                                             region #357.
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New nucleic acid sequences of the optineurin gene are useful to detect polymorphisms particularly single nucleotide polymorphisms in the optineurin promoter to diagnose, prognose and treat glaucoma and related disorders.

Claim 11; SEQ ID NO 359; 159pp; English.

Creat 20 but not more than 1500 consecutive nucleotides of the optineurin comprashly linked to a heterologous nucleic acid, a nucleic acid capable of detecting a single nucleotide polymorphism (SNP) in the optineurin promoter coperably linked to a heterologous nucleic acid, a nucleic acid capable of cetecting a single nucleotide polymorphism (SNP) in the optineurin promoter promoter, a host cell comprising the promoter operably linked to a compression of the optineurin gene, associated with a sample containing a SNP sequence variation in a sample containing a SNP sequence variation in a sample containing DNA, detecting a SNP sequence variation in a sample containing bus of visual field in a patient (or the severity corporatesistion primers that direct amplification reaction primers that direct amplification reaction primers that direct amplification a sample containing the DNA) and detecting a polymorphism cobatining a sample containing the DNA) and detecting a polymorphism cobatining a sample containing the DNA) and detecting a polymorphism of capable of detecting a SNP located within an optineurin promoter, and detecting the polymorphism). The invention is used to diagnose and corporate sequence is an optineurin promoter motif, repeat element or present sequence is an optineurin promoter motif, repeat element or containing the primer motif, repeat element or put attitute regulatory region.

Sequence 32 BP; 5 A; 12 C; 8 G; 7 T; 0 U; 0 Other;

Ś Matches Query Match Best Local S 860 AAGTGCTGGGATTACAGGCGTGAGCCACCACG 31 ; Similarity Conservative 3.1%; <u>.</u>. Score Pred. Mismatches 2 30.4; LL 30.5e+02; 891 1; 1, Length 32; 0, Gaps

0

RESULT 247

AAH91142/C

ID AAH91142 standard; DNA; 36 BP.

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AC AAH91142;

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32

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PXSXQXEXPXSXI
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                                                                                                                                                                             RESULT 248
ACC84462
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                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
Matches 32; Conserv
                                                                                                                                                                                                                                                                                                                                                      The present invention describes a method for detecting the presence of polymorphisms associated with inflammatory bowel diseases such as ulcerative colitis and Crohn's disease. The methods can be used to detect the presence of genetic polymorphisms associated with inflammatory bowel disease and correlating their occurrence with disease states. They may be used in this way for phenotypic correlations, forensics, paternity testing, medicine and genetic analysis. The present sequence is a polymorphic site described in the exemplification of the invention
WO2003008443-A2
                         Unidentified
                                                                                                                                                               ACC84462 standard;
                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Testing for the presence of polymorphisms associated with inflammatory bowel disease, using a hybridization assay.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Daly M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10-DEC-1999; 99US-0170257P.
10-APR-2000; 2000US-0196046P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  11-DEC-2000; 2000WO-US033632
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           14-JUN-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   chromosome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; inflammatory bowel disease; Crohn's disease; ulcerative colitis; single nucleotide polymorphism; SNP; chromosome 19p13; paternity test;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human inflammatory bowel disease associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             09-OCT-2001
                                                neural thread
                                                              Cytostatic;
                                                                                     NTP peptide
                                                                                                               28-AUG-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (ELLI-)
                                                                                                                                                                                                                                                       1032
                                                                                                                                                                                                                               36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WHITEHEAD INST BIOMEDICAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ELLIPSIS BIOTHERAPEUTICS CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hudson TJ,
                                                                                                                                                                                                                               AGCTGGGATTACAGGCANCTGCCACCACGCCCGGCT 1
                                                                                                                                                                                                                                                       AGCTGGGATTACGGGCACCTGCCACCACACCCCGCT 1067
                                                                                                                                                                                                                                                                                                                                 36 BP; 5 A; 10 C; 13 G; 7 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Page 48; 463pp; English
                                               Antibacterial; Immunosuppressive; Antiinflammatory; ad protein; NTP; tumour; ds.
                                                                                   encoding sequence
                                                                                                                                                                                                                                                                                 Conservative
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                                                                                                             (first
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /*tag= a
/note= "SNP, optionally G or T at this position"
                                                                                                                                                              DNA;
                                                                                                               entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  forensic test; gene
                                                                                                                                                                                                                                                                                          3.1%;
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                                                                                                                                                                 뫉
                                                                                                                                                                                                                                                                              0,
                                                                                                                                                                                                                                                                                            Score 30.2; DB 1;
Pred. No. 5.5e+02;
                                                                                     #9
                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Rioux J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  therapy;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   polymorphic site #217.
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                                                                                                                                                                                                                                                                                                        Length
                                                                                                                                                                                                                                                                                 Indels
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RESULT 249
ADE14029/c
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19-JUL-2001;
16-NOV-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               treating a condition in a patient requiring removal or destruction of cells, for treating a condition such as benign or malignant tumor, inflammatory disease, autoimmune disease and infectious disease. The peptide useful for treatment is derived from the amino acid sequence for a pancreatic thread protein. The peptide is conjugated, linked or bound to a molecule chosen from antibody or its fragment, antibody-like binding molecule, where the molecule has a higher affinity for binding to a tumor or other target than binding to other cells. Treatment using NTP peptides can remove benign tumors with less risk and fewer of the undesirable side effects of surgery. The present sequence is an NTP encoding sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   referred to as cell death peptide. Thought to be cyroscart, referred to as cell death peptide. Thought to be cyroscart, antibacterial, immunosuppressive and antiinflammatory. It is useful for antibacterial, immunosuppressive and antiinflammatory. It is useful for antibacterial, immunosuppressive and antiinflammatory.
                                                                                                                                                                                                                    SNP; glaucoma; progressive ocular hypertensive disorder; glaucoma related disorder; motif; repeat element; regulatory region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel neural thread protein peptide, referred as cell death peptide, useful for treating prostatic hyperplasia, psoriasis, eczema, dermatos atherosclerosis, cosmetic modification to skin, throat, mouth, muscle.
                                                                                                                                                                                                                                                                                                                           29-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 30 BP; 6 A; 9 C; 10 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention relates to a neural thread protein (N' referred to as cell death peptide. Thought to be cytostatic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               P-PSDB; ABR63257.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2003-247999/24.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19-JUL-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30-JAN-2003
                                               06-MAR-2002; 2002US-00091281
                                                                               06-MAR-2002; 2002US-00091281.
                                                                                                                 09-OCT-2003
                                                                                                                                                  US2003190617-A1
                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                      Human; optineurin; ds; ophthalmological; single nucleotide polymorphism;
                                                                                                                                                                                                                                                                                       Optineurin promoter motif, repeat element
                                                                                                                                                                                                                                                                                                                                                                                            ADE14029
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              \vdash
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PA;
                                                                                                                                                                                                                                                                                                                                                                                            standard; DNA; 32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Page 17; 77pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3.0%;
llarity 100.0%;
Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2001US-0306150P.
2001US-0306161P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2002WO-CA001105
                                                                                                                                                                                                                                                                                                                         (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 30;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  5e+02;
                                                                                                                                                                                                                                                                                     or regulatory region #138.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (NTP) peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ٥,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               dermatosis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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(SIEE/)

SI E. RAYMOND V.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CC heterologous sequence, diagnosing or prognosing glaucoma in a sample contained from a cell or bodily fluid (comprising detecting a polymorphism clin a promoter region of the optineurin gene, associated with a glaucoma clin a promoter region of the optineurin gene, associated with a glaucoma components, detecting a SNP sequence variation in a sample containing DNA, detecting the presence of an optineurin promoter sequence variation clin a sample containing DNA, determining the presence or increased containing the presence or increased containing the presence or increased containing in loss of visual field in a patient (or the severity containing the presence or increased containing in loss of visual field in a patient (or the severity contribution of glaucoma in a patient, comprising providing complification reaction primers that direct amplification of a selected containing a manufacture that direct amplification of a selected containing a the DNA) and detecting a polymorphism (comprising containing the DNA) and detecting a polymorphism (comprising containing a sample containing an unclaid and containing a sample containing and also to treat glaucoma related disorders. The consense and also to treat glaucoma related disorders. The containing an optineurin promoter motif, repeat element or containing promoter motif, repeat element or put at tive regulatory region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local S
Matches 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New nucleic acid sequences of the optineurin gene are useful to detect polymorphisms particularly single nucleotide polymorphisms in the optineurin promoter to diagnose, prognose and treat glaucoma and related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       operably linked to a heterologous nucleic acid, a nucleic acid capable detecting a single nucleotide polymorphism (SNP) in the optineurin promoter, a host cell comprising the promoter operably linked to a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                least 20 but not more than 1500 consecutive nucleotides of the optineurin promoter appearing as ADE13890. Also included are the optineurin promoter operably linked to a heterologous nucleic acid, a nucleic acid capable of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 11; SEQ ID NO 140; 159pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Raymond V, Morissette
                                                                                                                                                                                                                                                                          Enhancer element er-4 conserved basepair sequence.
                                                                                                                                                                                                                                                                                                               25-MAR-2003
25-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                AAQ73572 standard; DNA; 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (MORI/) MORISSETTE
                                                                                                                                                                                                              Enhancer element; carcinoma; tumor; cancer; SLPI gene; secretory leukoprotease-inhibitor gene; cytokeratin ge
                                                       misc_difference
                                                                                                           misc_difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         invention relates to an isolated nucleic acid (N1) comprising st 20 but not more than 1500 consecutive nucleotides of the opi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2003-864168/80
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAGTGCTGGGATTACAGGCGTGAGCCACCAC 890
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP; 5 A; 11 C; 9 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                   (revised)
                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                               Location/Qualifiers
                  /*tag=
/label=
                                                                          label=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 96.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3.0%;
                b
pyrimidine
                                                                        purine
                                                                                                                                                                                                                                                                                                                                                                                                                    BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 29.4;
Pred. No. 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    .5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 32;
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                                                                                                                                                                                                                gene-8;
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Best Local S
Matches 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This enhancer element is part of a DNA construct used for treating human carcinoma which contains a cancer therapeutic protein under the control of a promoter and 3 enhancer sequences in a specific 5'-3' order. This enhancer element is derived from the flanking region of the human epithelial cell cytokeratin-8 gene. (Updated on 25-MAR-2003 to correct PN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1; Fig 6; 54pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA construct for treating human carcinoma - therapeutic gene under the control of a promother property of the control of the control of the property of the control of th
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                                                                                                                                                                               07-MAY-1998;
03-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Lesch-Nyhan syndrome; muscular dystrophy; Wiskott-Aldrich syndrome; Fabrys disease; familial hypercholesterolemia; hereditary spherocytosis; polycystic kidney disease; von Willebrands disease; forensic; human; tuberous sclerosis; hereditary hemorrhagica telanglectasia; familial colonic polyposis; osteogenesis imperfecta; porphyria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAA04659;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAA04659 standard; DNA; 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    29-SEP-1994
                                                                                                                                                                                                                                                                                                                                                            EP955382-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Polymorphic fragment of hypertension associated
                                             Fan JB,
                                                                                                                                                                                                                                                        07-MAY-1999;
                                                                                                                                                                                                                                                                                                           10-NOV-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ehlers-Danlos
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Polymorphism;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (UABR-) UAB RES FOUND
                                                                                                                                                                                                                                                                                                                                                                                                               Omo
                                                                                                 (AFFY-) AFFYMETRIX INC.
(UYCA-) UNIV CASE WESTI
                                                                                                                                                                                                                                                                                                                                                                                                            sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CTCAGCCTCCCAAAGTGCTGGGATTACAGGC 408
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CTCAGCCTCCCAANGTGCTGGGANTACAGGC 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          31 BP; 6 A; 10 C; 8 G; 5 T; 0 U; 2 Other;
                                             Chakravarti A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sorscher
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                   CASE WESTERN RESERVE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       hypertension; agammaglobulinemia; diabetes insipidus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              93US-00035435
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                                                                                                                                                                            98US-0084641P.
99US-00304232.
                                                                                                                                                                                                                                                        99EP-00250150
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93.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             띰
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  88.
                                                   Haluska MK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 29; I
Pred. No. 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1;
5.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           oma - includes a cancer-
promoter and a gp. of e
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 31
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
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WPI; 2000-107928/10

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RESULT 252
AAM379
AAM379
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AAM379
AAM
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 kidney disease, hereditary spherocytosis, von Willebrands disease, tuberous sclerosis, hereditary hemorrhagica telangiectasia, familial colonic polyposis, Ehlers-Danlos syndrome, osteogenesis imperfecta, a acute intermittent porphyria. The polymorphic forms can also be used forensics to identify individuals
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            regimes customized to underlying abnormalities. The polymorphisms can be used for association studies for hypertension, and in hypertension diagnostic assays. Where the polymorphisms have strong correlation with hypertension, within a gene, they are likely to have a causative role in hypertension. This information can be used to find the precise role of a polymorphism in the disease, and this can be used to identify potential drugs which combat the disease, and this can be used to identify potential drugs which combat the disease. The polymorphisms can be tested for association with other diseases e.g. agammaglobulinemia, diabetes insipidus, Lesch-Nyhan syndrome, muscular dystrophy, Wiskott-Aldrich syndrome, Fabrys disease, familial hypercholesterolemia, polycystic video. Fabrys disease, familial hypercholesterolemia, polycystic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis; inflammation; forensic investigation; paternity analysis; PCR primer; ss.
                               New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nuc
                                                                                                                                      WPI; 2001-290930/30
                                                                                                                                                                                                 Picoult-Newburg
                                                                                                                                                                                                                                                                                                                                                                                                     13-OCT-2000; 2000WO-US028436.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO200129262-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAH37977;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAH37977 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention provides polymorphic fragments of genes associated with hypertension. The nucleic acids including the polymorphic sites can be used as probes or primers for expressing variant proteins. Detection of the polymorphisms is useful in designing prophylactic and therapeutic the polymorphisms is useful in designing prophylactic and therapeutic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 1; Page 43;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              hypertension
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                                                                                                                                                                                                                                                                                                                                    15-OCT-1999;
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                                                                                                                                                                                                                                                                     ORCHID
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CCCAGGCTGGAGTGCAGTGGCGCAATCT 670
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP; 5 A; 8 C; 10 G; 5 T; 0 U;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                     BIOSCIENCES INC
                                                                                                                                                                                                                                                                                                                                    99US-0160096P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PCR primer SEQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 27.6; DB 1; Length 29; Pred. No. 6.3e+02; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ID 773.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
                                  nucleic
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in
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Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide primer extension (SNPB) primers, and the sequences of regions flanking sites of single nucleotide polymorphisms SNPs. The present invention includes kits for determining the presence or absence of a SNP, using the disponucleotides of the invention. The PCR primers are used to amplify a SNP flanking sequence, the SNPE primer is used as a genotyping primer. The oligonucleotides are useful for genotyping a mucleic acid sample by performing a single-nucleotide primer extension reaction. The oligonucleotides are useful for determining the presence, absence or identity of a SNP and for genotyping nucleic acid samples, for e.g. to assess by association analysis the genotype of an individual or group of individuals, having a pathological phenotypic trait suspected of being caused by one or more SNPs. Phenotypic traits include diseases e.g. agammaglobulinaemia, diabetes insipidus, Lesch-Nyhan syndrome, muscular agammaglobulinaemia, diabetes insipidus, Lesch-Nyhan syndrome, muscular
                                                                               inflammation, cancer, nervous system diseases and infection by pathogenic microorganism. The method is also useful in forensic investigations and paternity analysis. The present sequence represents a PCR primer specific
                                                                                                                                                                     agammaglobulinaemia, diabetes insipidus, Lesch-Nyhan syndrome, musculidystrophy, familial hypercholesterolaemia, polycystic kidney disease, osteogenesis imperfecta and acute intermitent porphyria. Phenotypic traits also include symptoms of or susceptibility to multifactorial disease of which a component is or may be genetic such as autoimmune diseases, including, rheumatoid arthritis, multiple sclerosis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1; Page 53; 83pp; English
Sequence
                                                             for a human
      29
   B₽;
                                                             SNP
      12
                                                          containing DNA sequence
A; 7 C; 3 G; 7 T; 0 U;
      0 Other;
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Matches Query Match Local 774 29 Similarity GTATTTTAGTAGAGATGGGTTTCACCAT 1 GTATTTTAGTAGAGATGGGGTTCACCAT 802 Conservative 96.6%; 0; Score 27.4; DB 1; Pred. No. 6.4e+02; Mismatches Length Indels 29; 0, Gaps

0

밁 8

RESULT 253
AAXO64467
ID AAXO64
XX AAXO64
XX AAXO64
XX J-MAH
XX POlyme
KW ACC
XX ACC
PL AAXO64
XX ACC
XX ACC
PL AAXO64
XX ACC
XX ACC Human biallelic polymorphic DNA fragment SGC34924. 31-MAR-1999 AAX06467 standard; DNA; 31 BP (first entry)

Polymorphism; biallelic; paternity testing; forensic; genetic mapping; phenotypic typing; medicament; disease; marker; human; ss.

Homo sapiens.

WO9858529-A2.

30-DEC-1998

22-JUN-1998; 98WO-US012930

97US-0050594P

(AFFY-) AFFYMETRIX INC.

Lipshutz 핂, Chee M, Fan Berno

WPI; 1999-080963/07.

New nucleic acid segments containing polymorphic sites - used for, edetecting a disease phenotype, in forensics, paternity testing or generapping of phenotypic traits. genetic

Claim 1; Page 29; 61pp; English

The

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RESULT 254
AAQ27389
ID AAQ273
XX AAQ273
AC AAQ273
XX DT 25-MAR
DT 27-JAN
XX Polymm
XX Polymm
XX Polymm
XX WO921:
XX WO921:
XX WO921:
XX Uittes
PR 24-JAN
XX Uittes
PR 25-JAN
XX Uittes
PR WPI;
XX Uittes
PR WPI;
XX Detect
PT hybrid
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    application programme and including records, each record comprising information identifying a polymorphism shown in the above sequences. products and methods can be used for analysing polymorphic sites in individuals for testing for the presence of a disease phenotype or in forensics, paternity testing or genetic mapping of phenotypic traits.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAQ27389 standard; DNA; 32
                        primer PDJ33A is one of several primers which are preferred for use in amplifying inter-Alu regions of DNA. The amplified fragments are then subjected to 2-D electrophoresis on the basis of length and differences in base sequence. The resulting separation pattern is transferred to a filter for screening with a probe. The method can be used to detect genetic variation. See also AAQ27390-Q27404 and AAQ33141-Q33144. (Update on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    25-MAR-2003
27-JAN-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAQ27389;
                                                                                                                                                                                                                                                    Detection of genetic variation by 2-D electrophoresis of fragments - and hybridisation with labelled probes, carried out on fragments consisting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           24-JAN-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO9213101-A1
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                                                                                                                                                                                                                                                                                                                                                               Uitterlinden AG,
                                                                                                                                                                                                                                                                                                                                                                                                         (INGE-) INGENY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        06-AUG-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Polymerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Inter-Alu specific primer PDJ33A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local
                                                                                                                                                                                                                              inter-repeat sequences generated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        y can also be used for the production of polypeptides expressed by lant genes and for the production of transgenic animals. The nucleic segments can also be used in the manufacture of medicaments for the
                                                                                                                                                                                                                                                                                                                       1992-284683/34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               863
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    or prophylaxis of diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    chain reaction; PCR; repetitive element;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP; 8 A; 9 C; 8 G; 5 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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(first en
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                                                                                                                                                                                          31pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    entry)
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Pred.
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No. 6.
                                                                                                                                                                                                                                    carried by PCR.
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(Updated

detecting the polymorphism). The invention is used to diagnose and prognose glaucoma and also to treat glaucoma related disorders. The

motit

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RESULT 255
ADE14206/c
ID ADE142
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S
                                      The invention relates to an isolated nucleic acid (N1) comprising at CC least 20 but not more than 1500 consecutive nucleotides of the optineurin CC promoter appearing as ADE13890. Also included are the optineurin promoter operably linked to a heterologous nucleic acid, a nucleic acid capable of CC detecting a single nucleotide polymorphism (SNP) in the optineurin CC promoter, a host cell comprising the promoter operably linked to a CC heterologous sequence, diagnosing or prognosing glaucoma in a sample CC obtained from a cell or bodily fluid (comprising detecting a polymorphism CC in a promoter region of the optineurin gene, associated with a glaucoma CC phenotype), detecting a SNP sequence variation in a sample containing DNA, determining the presence or increased CC in a sample containing DNA, determining the presence or increased CC disorder resulting in loss of visual field in a patient (or the severity or progression of glaucoma or to a progressive ocular hypertensive CC amplification reaction primers that direct amplification of a selected containing a sample containing the variation within the optineurin obtaining a sample containing the variation within the optineurin compaties of detecting a SNP located within an optineurin promoter, and comprising of detecting a SNP located within an optineurin promoter, and detecting a providing a nucleic acid capable of detecting a SNP located within an optineurin promoter, and companies and the providing and companies and the providing and comprising of the providing and providing a nucleic acid capable of detecting a SNP located within an optineurin promoter, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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Best Local &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 32 BP; 7 A; 9 C; 10 G; 6 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New nucleic acid sequences of the optineurin gene are useful to detect polymorphisms particularly single nucleotide polymorphisms in the optineurin promoter to diagnose, prognose and treat glaucoma and related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SNP; glaucoma; progressive glaucoma related disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Optineurin promoter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADE14206 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; optineurin; ds; ophthalmological; single nucleotide polymorphism;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        29-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADE14206;
                                                                                                                                                                                                                                                                                                                                                                                                                   Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Raymond V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    06-MAR-2002; 2002US-00091281
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     06-MAR-2002; 2002US-00091281
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (SIEE/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SI E.
RAYMOND V.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MORISSETTE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TCAGCCTCCCAAAGTGCTGGGATTACAGG
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                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID NO 317;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Morissette
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                disorder; motif;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       motif,
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                                                                                                                                                                                                                                                                                                                                                                                                                     159pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ocular
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     repeat
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 27.4;
Pred. No. 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        407
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     element;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     regulatory region
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RESULT 256
ANI62688/c
ID ANI62688;
XX IJ-OCT-200
XX Human; brea
XX IJ-JAN-200
PR 10-ANR-200
PR 24-FEB-200
PR 24-FEB-200
PR 11-ANR-200
PR 11-ANR-200
PR 11-ANR-200
PR 11-ANG-200
PR 12-ANG-200
PR 22-ANG-200
PR 23-ANG-200
PR 23-ANG-200
PR 11-SEP-200
PR 01-SEP-200
PR 01-SEP-200
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Best Local S
Matches 28
28-JUN-2000

30-JUN-2000

07-JUL-2000

11-JUL-2000

11-JUL-2000

14-JUL-2000

14-JUL-2000

14-AUG-2000

15-AUG-2000

16-AUG-2000

17-AUG-2000

18-AUG-2000

19-AUG-2000

20-AUG-2000

20-AU
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07-JUN-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; breast antigen; ovarian antigen; cancer; metastasis; gene therapy; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human breast
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19-OCT-2001
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1 Similarity 96.6%;
28; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 standard; DNA; 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         32 BP; 7
2000US-0179065P
2000US-0184664P
2000US-0184664P
2000US-0184664P
2000US-0199076P
2000US-0199076P
2000US-029467P
2000US-0216487P
2000US-0216487P
2000US-0216487P
2000US-0216487P
2000US-0217487P
2000US-0218290
2000US-02245189
2000US-0224518P
2000US-02255267P
2000US-0225266P
2000US-0225266P
2000US-0225266P
2000US-02252686P
2000US-02252686P
2000US-0225759P
2000US-0225758P
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No
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     01-SEP-2000
05-SEP-2000
06-SEP-2000
06-SEP-2000
08-SEP-2000
08-SEP-2000
08-SEP-2000
08-SEP-2000
08-SEP-2000
11-SEP-2000
12-SEP-2000
12-SEP-2000
12-SEP-2000
13-OCT-2000
11-NOV-2000
 2000US-0229345P
2000US-023943P
2000US-023143P
2000US-0231243P
2000US-0231243P
2000US-0231414P
2000US-0231414P
2000US-0231414P
2000US-0231414P
2000US-0231414P
2000US-023239P
2000US-023239P
2000US-023239P
2000US-023239P
2000US-023363P
2000US-023499P
2000US-0241789P
2000US-0241789P
2000US-0241789P
2000US-024676P
2000US-024677P
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2001WO-US001339

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RESULT 257
AAL06807/c
X B X S X W W X B X D X A X I I
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Best Local Similarity
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17-NOV-2000;
17-NOV-2000;
01-DEC-2000;
01-DEC-2000;
05-DEC-2000;
05-DEC-2000;
06-DEC-2000;
08-DEC-2000;
                                                                                                                                                                                                                                                                                                                                                                                             The present invention provides the protein and coding sequences of a number of ovarian and breast antigens. These are shown in AAI62467-AAI62572 and AAM42240-AAM42345. The sequences can be used in the diagnosis, prevention and treatment of breast and ovarian cancers, and their metastasses. The present sequence is a genomic sequence of the invention. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17-NOV-2000;
17-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         17-NOV-2000;
17-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New isolated nucleic acids and polypeptides, useful for diagnosing treating and/or preventing human diseases and disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; SEQ ID NO
                                                                                                                                     21-NOV-2001
                                                                                                                                                                                           AAL06807
                                                                                                                                                                                                                                                                                                                                                                         Sequence 33 BP;
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                                                                                                         Human
                                       Homo sapiens
                                                                 cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   7-NOV-2000;
7-NOV-2000;
7-NOV-2000;
7-NOV-2000;
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                                                                                                                                                                                                                                                                               932 TCACTCTGTTACCCAGGCTGGAGTGCAATGGC 963
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                                                                                                        reproductive system related antigen DNA SEQ ID NO:
                                                                                                                                                                                                                                                                                                                     29;
                                                                               reproductive system related antigen; reproductive system
                                                                 gene
                                                                                                                                                                                            standard; DNA;
                                                                                                                                                                                                                                                               TCGCTCTGTTGCCCAGGCTGGAGTGCAGTGGC 2
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2000US-024921AP.
2000US-024921AP.
2000US-024924AP.
2000US-024924AP.
2000US-024926AP.
2000US-024926AP.
2000US-024926AP.
2000US-024929P.
2000US-024929P.
2000US-025931AP.
2000US-025931AP.
2000US-0251988P.
2000US-0251866P.
2000US-0251866P.
2000US-0251866P.
2000US-0251866P.
2000US-0251866P.
2000US-0251869P.
2000US-0251989P.
2000US-0251989P.
2000US-0251989P.
2000US-0251999P.
2000US-025199P.
2000US-025199P.
                                                                                                                                                                                                                                                                                                                      Conservative
                                                                  therapy;
                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                          8 A;
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                                                                                                                                                                                                                                                                                                                                                                          12 C; 9 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              338; 520pp + Sequence Listing; English.
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Pred. No. 7.2e+02;
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                                                                                  disorder;
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  28-UNN-2000;
28-UNN-2000;
30-UNN-2000;
31-UNN-2000;
11-UNL-2000;
11-UUL-2000;
11-UU
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24-FEB-2000;
02-MAR-2000;
16-MAR-2000;
17-MAR-2000;
18-APR-2000;
19-MAY-2000;
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2000US-0227182P
2000US-0228924P
2000US-0229344P
2000US-0229344P
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2000US-0229343P
2000US-0230437P
2000US-0231244P
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2000US-0235344P
2000US-023534PP
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2000US-0179065P 2000US-0184664P 2000US-0184664P 2000US-0184664P 2000US-0184664P 2000US-0199076P 2000US-0199076P 2000US-029467P 2000US-021648PP 2000US-021648PP 2000US-021748PP 2000US-021748PP 2000US-021749PP 2000US-0224518PP 2000US-0224519P 2000US-0225714PP 2000US-0225714PPP 2000US-0225714PPP 2000US-0225714PPP 2000US-0225714PPPPPPPPPP

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WPI; 2001-465570/50
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17-NOV-2000;
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                     Barash SC,
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2000US-0246528P
2000US-0246528P
2000US-0246532P
2000US-0246609P
2000US-0246610P
2000US-0246611P
2000US-0246611P
2000US-0249207P
2000US-0249207P
2000US-0249211P
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2000US-0251479P
2000US-0251868P
2000US-0251868P
2000US-0251869P
2000US-0251989P
2000US-0251990P
2000US-0251990P
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2000US-0246478P.
2000US-0246523P.
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2000US-
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2000US-0237038P.
                                             GENOME
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RESULT 258
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Matches
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                       The invention relates to a novel HOMO glandulae mammaria susceptibility gene encoding protein 12.32. The protein can be expressed by standard recombinant methodology. The protein and encoding polynucleotides are used in diagnosis and treatment of embryo peroplasia and cancer. The present sequence represents the HOMO glandulae mammaria susceptibility protein 12.32 cDNA cloning PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention provides the protein and coding sequences of a number of human reproductive system related antigens. These can be used in the prevention and treatment of reproductive system disorders, including cancer. The present sequence is a genomic sequence encoding a protein of the invention
Sequence 33
                                                                                                                                              A novel HOMO glandulae mammaria susceptibility gene encoding protein 12.32, and its polynucleotide, useful in the diagnosis and treatment embryo peroplasia and cancer.
                                                                                                                                                                                                                                Mao
                                                                                                                                                                                                                                                                                                                                                                                                                      Glandulae mammaria susceptibility protein; 12.32; cytostatic; human; gene therapy; cancer; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Isolated nucleic acid molecule encoding a reproductive used in preventing, treating or ameliorating a medical
                                                                                                                      Example
                                                                                                                                                                                                      WPI; 2002-281239/32.
                                                                                                                                                                                                                                                                                    21-JUN-2000; 2000CN-00116681.
                                                                                                                                                                                                                                                                                                                                                                  WO200220576-A1.
                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
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                                                                                                                                                                                                                                                          (BIOW-) BIOWINDOW GENE DEV INC SHANGHAI.
                                                                                                                                                                                                                                                                                                             19-JUN-2001; 2001WO-CN001005.
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                                                                                                                                                                                                                                                                                                                                                                                                                     therapy; cancer;
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                                                                                                                     4; Page 13; 38pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TCGCTCTGTTGCCCAGGCTGGAGTGCAGTGGC
 BP; 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              mammaria susceptibility gene
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9 C; 8 G; 10 T; 0
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                                                                                                                                                                                                                                                                                                                                                                                                                     PCR primer;
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Pred. No. 7.
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924 ATGGAATCTCACTCTGTTACCCAGGCTGGAGT

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Query Match Best Local S Matches 29

Similarity 29; Conserv

Conservative

2.7%;

Score 27.2; DB 1 Pred. No. 7.2e+02 0; Mismatches

DB 1;

Length 33; Indels

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ATGGATCCTCATTCTGTTACCCAGGCTGGAGT 33

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RESULT 259
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                                                                 RESULT 260
AAA04371
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                                                                                                                                                                             Matches
                                                                                                                                                                                             Best
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                                                                                                                                                                                                                                                                     The present invention relates to a neural thread protein (NTP) peptide referred to as cell death peptide. Thought to be cytostatic, antibacterial, immunosuppressive and antiinflammatory. It is useful for treating a condition in a patient requiring removal or destruction of cells, for treating a condition such as benign or malignant tumor, inflammatory disease, autoimmune disease and infectious disease. The peptide useful for treatment is derived from the amino acid sequence for a pancreatic thread protein. The peptide is conjugated, linked or bound to a molecule chosen from antibody or its fragment, antibody-like binding molecule, where the molecule has a higher affinity for binding to a tumor or other target than binding to other cells. Treatment using NTP peptides can remove benign tumors with less risk and fewer of the undesirable side
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19-JUL-2001; 2001US-0306150P.
19-JUL-2001; 2001US-0306161P.
16-NOV-2001; 2001US-0331477P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cytostatic; Antibacterial; Immunosuppressive; Antiinflammatory; neural thread protein; NTP; tumour; ds.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel neural thread protein peptide, referred as cell death peptide, useful for treating prostatic hyperplasia, psoriasis, eczema, dermatosis, atherosclerosis, cosmetic modification to skin, throat, mouth, muscle.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               30-JAN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NTP peptide encoding sequence
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                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 17; 77pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19-JUL-2002; 2002WO-CA001105
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                                                     AAA04371
                                                                                                                                                                                           Local
                                                                                                                                                  1017
                                                                                                                                                                                                                                                 remove benign tumors with 1908 and NTP encoding sequence
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                                                                                                                                                                             Similarity
27; Conser
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                                                     standard;
                                                                                                                                             CTCAGCCTCCCAAGCAGCTGGGATTAC 1043
                                                                                                                          CTCAGCCTCCCAAGCAGCTGGGATTAC
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Pred. No.
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                                                                                                                                                                              Mismatches
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22-MAY-2000

(first entry)

AAA04506
ID AAA(
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AC AAA(
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AC AAA(
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AT 22-N
XX

22-MAY-2000 AAA04506;

(first entry

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RESULT 261

AAA04506 standard; DNA; 29 BP

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The invention provides polymorphic fragments of genes associated with CC hypertension. The nucleic acids including the polymorphic sites can be CC used as probes or primers for expressing variant proteins. Detection of CC the polymorphisms is useful in designing prophylactic and therapeutic CC regimes customized to underlying abnormalities. The polymorphisms can be CC used for association studies for hypertension, and in hypertension of CC diagnostic assays. Where the polymorphisms have strong correlation with CC hypertension. This information can be used to find the precise role in CC hypertension. This information can be used to find the precise role of a CC polymorphism in the disease. The polymorphisms can be tested for CC drugs which combat the disease. The polymorphisms can be tested for CC association with other diseases e.g. agammaglobulinemia, diabetes CC insipidus, Lesch-Nyhan syndrome, muscular dystrophy, wiskott-Aldrich CC syndrome, Fabrys disease, familial hypercholesterolemia, polycystic C tuberous sclerosis, hereditary spherocytosis, von Willebrands disease, familial cc colonic polyposis, Ehlers-Danlos syndrome, osteogenesis imperfecta, and CC acute intermittent porphyria. The polymorphic forms can also be used in CC forensics to identify individuals
                                                                                                                          Matches
                                                                                                                                                 Query Match
Best Local S
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Losch-Nyhan syndrome; muscular dystrophy; Wiskott-Aldrich syndrome;
Fabrys disease; familial hypercholesterolemia; hereditary spherocytosis;
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03-MAY-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 1; Page 34; 53pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10-NOV-1999.
                                                                                                                                                                                                                                             Sequence 29 BP; 6 A; 8 C; 9 G; 5 T; 0 U; 1 Other;
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(UYCA-) UNIV CASE WESTERN RESERVE.
                             856 CCCAAAGTGCTGGGATTACAGGCGTGAGC 884
                                                                                                                          27;
μ
                                                                                                                                                    Similarity
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CCCAAAGTGCTGGGRTTACAGGCCTGAGC
                                                                                                                          Conservative
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99US-00304232.
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                                                                                                                                                    2.7%;
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                                                                                                                                                    Score
Pred.
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                                                                                                                                                              No.
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                                                                                                                                                                                        DB 1;
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                                                                                                                 RESULT 262
AAA04303
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                                                                                                                                                                                                                                                                                                                                                 regimes customized to underlying abhormalities. The polymorphisms can be CC used for association studies for hypertension, and in hypertension with diagnostic assays. Where the polymorphisms have strong correlation with hypertension, within a gene, they are likely to have a causative role in hypertension. This information can be used to find the precise role of a golymorphism in the disease, and this can be used to identify potential CC drugs which combat the disease. The polymorphisms can be tested for association with other diseases e.g. agammaglobulinemia, diabetes insipidus, Lesch-Nyhan syndrome, muscular dystrophy, Wiskott-Aldrich cyndrome, Fabrys disease, familial hypercholesterolemia, polycystic kidney disease, hereditary spherocytosis, von Willebrands disease, tuberous sclerosis, hereditary spherocytosis, von Willebrands disease, colonic polyposis, Ehlers-Danlos syndrome, osteogenesis imperfecta, and corrensics to identify individuals
                                                                                                                                                                                                                                                        Matches
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Best Local (
Polymorphic fragment of hypertension associated gene GLUT4.
                                   22-MAY-2000
                                                                    AAA04303;
                                                                                                   AAA04303 standard; DNA;
                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention provides polymorphic fragments of genes associated with hypertension. The nucleic acids including the polymorphic sites can be used as probes or primers for expressing variant proteins. Detection of the polymorphisms is useful in designing prophylactic and therapeutic the polymorphisms is useful in designing prophylactic and therapeutic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim
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03-MAY-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Lesch-Nyhan syndrome; muscular dystrophy; Wiskott-Aldrich syndrome; Fabrys disease; familial hypercholesterolemia; hereditary spherocytosis; polycystic kidney disease; von Willebrands disease; forensic; human; tuberous sclerosis; hereditary hemorrhagica telangiectasia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Polymorphism;
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                                                                                                                                                                                                         GCTGGGATTACAGGCGTGAGCCACCACGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Page
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                                                                                                                                                                                    GCTGGGATTACAGGYGTGAGCCACCGCGC
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                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                       BP; 5 A;
                               (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        syndrome; ss.
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99US-00304232.
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                                                                                                                                                                                                                                                                   2.7%;
                                                                                                                                                                                                                                                                                                                       8 C; 11 G; 4 T; 0 U; 1 Other;
                                                                                                   29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                English.
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                                                                                                                                                                                                                                                     1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     agammaglobulinemia;
                                                                                                                                                                                                                                                                    Score 27;
Pred. No.
                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                     892
                                                                                                                                                                                    29
                                                                                                                                                                                                                                                                                    DB 1;
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                                                                                                                                                                                                                                                                                    Length 29;
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AAA04500
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AC AAAC
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DT 22-N
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DE Poly
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                                                                                                                                                                                                                                                                                                                                                                                   CC typertension. The nucleic acids including the polymorphic sites can be cused as probes or primers for expressing variant proteins. Detection of the polymorphisms is useful in designing prophylactic and therapeutic regimes customized to underlying abnormalities. The polymorphisms can be used for association studies for hypertension, and in hypertension with CC diagnostic assays. Where the polymorphisms have strong correlation with CC hypertension, within a gene, they are likely to have a causative role in the polymorphism in the disease, and this can be used to identify potential CC drugs which combat the disease. The polymorphisms can be tested for association with other diseases e.g. agammaglobulinemia, diabetes clinsipidus, Lesch-Nyhan syndrome, muscular dystrophy, Wiskott-Aldrich CC inspiritus, Lesch-Nyhan syndrome, muscular dystrophy, Wiskott-Aldrich CC syndrome, Fabrys disease, familial hypercholesterolemia, polycystic kidney disease, hereditary spherocytosis, von Willebrands disease, tuberous sclerosis, hereditary spherocytosis, von Willebrands disease, colonic polyposis, Ehlers-Danlos syndrome, osteogenesis imperfecta, and CC colonic polyposis, Ehlers-Danlos syndrome, osteogenesis imperfecta, and CC forensics to identify individuals
                                                                                                                                                                                                                                                                                  Matches
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Polymorphism; hypertension; agammaglobulinemia; diabetes insipidus;
Lesch-Nyhan syndrome; muscular dystrophy; Wiskott-Aldrich syndrome;
Fabrys disease; familial hypercholesterolemia; hereditary spherocytosis;
                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention provides polymorphic fragments of genes associated with hypertension. The nucleic acids including the polymorphic sites can be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2000-107928/10
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tuberous sclerosis; hereditary hemorrhagica telangiectasia;
familial colonic polyposis; osteogenesis imperfecta; porphyria;
                                                      22-MAY-2000 (first entry)
                                                                                                                         AAA04500 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1; Page 32; 53pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel nucleic acids containing polymorphisms used
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03-MAY-1999;
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                                                                                                                                                                                                                                                                                Local Similarity nes 27; Conserv
                                                                                                                                                                                                                                                862 GTGCTGGGATTACAGGCGTGAGCCACCAC
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                                                                                                                                                                                                               GTGCTGGGATTACARGCGTGAGCCACCGC
                                                                                                                                                                                                                                                                                                                                                    BP; 5 A; 8 C;
                                                                                                                                                                                                                                                                                  Conservative
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99US-00304232.
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                                                                                                                                                                                                                                                                                               2.7%;
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                                                                                                                                                                                                                                                                                Score 27; DB
Pred. No. 6.7e
1; Mismatches
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                                                                                                                                                                                                                                                                                                   6.7e+02;
                                                                                                                                                                                                                                                                                                                                                       U; 1 Other;
                                                                                                                                                                                                                                                                                                                   Length 29;
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                                                                                                                                                                                                                                                                                    Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               the diagnosis
                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                  Gaps
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Polymorphic fragment of hypertension associated gene PGIS

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RESULT 264
AAA0396
ID AAA039
XX AAA039
XX AAA039
XX DT 22-MA1
XX POlyma
XX POlyma
XX POlyma
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                                                                                                                                                                                                                                                                                                               Matches
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     hypertension
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel nucleic acids containing polymorphisms used in the diagnosis of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2000-107928/10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        07-MAY-1998;
03-MAY-1999;
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                                                                                                                                                                                                                                                                                                                                                                                      Sequence 29
                                                                                                                                                                                                                                                                                                                                                                                                                                         colonic polyposis, Ehlers-Danlos syndrome, osteogenesis imperfecta, a acute intermittent porphyria. The polymorphic forms can also be used
Polymorphism;
                                                                          22-MAY-2000
                                                                                                                                               AAA03996 standard; DNA; 29 BP
                                                                                                                                                                                                                                                                                                                                                                                                                          forensics to identify individuals
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                                                                                                                                                                                                                                                                            713
                                                                                                                                                                                                                                                                                                               27;
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                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                             polyposis,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Page 38; 53pp;
                                                                                                                                                                                                                                                         CTGCCCCAGCCTCCTGAGTAGCTGGGACT 741
                                                                                                                                                                                                                                        CTGCCTCAGCCTCCYGAGTAGCTGGGACT 29
                                                                                                                                                                                                                                                                                                                                                                                      BP; 4 A; 10 C; 8 G; 6 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                               Conservative
                                                                          (first entry)
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hypertension; agammaglobulinemia; diabetes insipidus;
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99US-00304232.
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93.1%;
                                      of hypertension associated gene APOC4.
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1; Mismatches
                                                                                                                                                                                                                                                                                                                                    Score 27;
Pred. No.
                                                                                                                                                                                                                                                                                                                                    6.7e+02;
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AAA04505
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AC AAAC
XX 22-P
XX Pol;
XX Pol;
XX Pol;
XX Les

(first entry)

AAA04505 standard; DNA; 29

ВP

0

Polymorphic 22-MAY-2000

Polymorphism; hypertension; agammaglobulinemia; diabetes insipidus; Lesch-Nyhan syndrome; muscular dystrophy; Wiskott-Aldrich syndrome;

fragment of hypertension associated gene PGIS

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RESULT 265
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                                                                                                                                                                                    Query Match
Best Local S
Matches 27
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03-MAY-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                hypertension.
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                                                                                                                                                                                                                                                                                             Sequence 29
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                                                                                                        849 TCGGCCTCCCAAAGTGCTGGGATTACAGG 877
                                                                                                                                                                                       27;
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                                                                                                                                                                                                                   Similarity
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                                                                               TTGGCCTCCCAAAGYGCTGGGATTACAGG
                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                             BP; 6 A; 7 C; 9 G; 6 T; 0 U; 1 Other;
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99US-00304232.
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                                                                                                                                                                                                                   93.1%;
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                                                                                                                                                                                                                Score 27; DB 1;
Pred. No. 6.7e+0;
                                                                                                                                                                                          Mismatches
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                                                                                                                                                                                                                   .7e+02
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                                                                                                                                                                                                                                         Length 29
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RESULT 266
AAQ73570
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention provides polymorphic fragments of genes associated with the prevention. The nucleic acids including the polymorphic sites can be used as probes or primers for expressing variant proteins. Detection of the polymorphisms is useful in designing prophylactic and therapeutic regimes customized to underlying abnormalities. The polymorphisms can be used for association studies for hypertension, and in hypertension with hypertension, within a gene, they are likely to have a causative role in hypertension, within a gene, they are likely to have a causative role in hypertension. This information can be used to find the precise role of a polymorphism in the disease, and this can be used to identify potential drugs which combat the disease. The polymorphisms can be tested for a sesociation with other diseases e.g. agammaglobulinemia, diabetes insipidus, Lesch-wyhan syndrome, muscular dystrophy, wiskott-Aldrich syndrome, Fabrys disease, familial hypercholesterolemia, polycystic kidney disease, hereditary spherocyposis, von Willebrands disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local S
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Enhancer element; carcinoma; tumor; cancer; SLPI gene;
secretory leukoprotease-inhibitor gene; cytokeratin gene-8;
                                                                                             Enhancer element er-3 conserved basepair sequence
                                                                                                                                                         25-MAR-2003
25-JUN-1995
                                                                                                                                                                                                                                                    AAQ73570;
                                                                                                                                                                                                                                                                                                         AAQ73570 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                tuberous sclerosis, hereditary hemorrhagica telangiectasia, familial colonic polyposis, Ehlers-banlos syndrome, osteogenesis imperfecta, acute intermittent porphyria. The polymorphic forms can also be used forensics to identify individuals
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1; Page 38; 53pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2000-107928/10
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03-MAY-1999;
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UNIV CASE WESTERN RESERVE.
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(first en
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99US-00304232.
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                                                                                                                                                         entry)
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Pred. No.
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L; Mismatches
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Best Local
24-FEB-2000;
02-MAR-2000;
16-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This enhancer element is part of a DNA construct used for treating human carcinoma which contains a cancer therapeutic protein under the control of a promoter and 3 enhancer sequences in a specific 5'-3' order. This enhancer element is derived from the flanking region of the human epithelial cell cytokeratin-8 gene. (Updated on 25-MAR-2003 to correct Plepithelial cell cytokeratin-8 gene. (Updated on 25-MAR-2003)
                                                                  31-JAN-2000;
04-FEB-2000;
                                                                                                                                                                                                                                                                                                                                               Human; digestive system antigen; gene therapy; cancer; ulcerative colitis; infection; Hirschsprung's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 32
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                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                     Human digestive system antigen genomic sequence SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  05-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA construct for treating human carcinoma - therapeutic gene under the control of a promoter than the control of the promoter of the control of the promoter of the control of the contro
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                                                                                                                                       17-JAN-2001;
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2000US-0184664P.
2000US-0186350P.
2000US-0189874P.
                                                                  2000US-0179065P.
2000US-0180628P.
                                                                                                                                          2001WO-US001324.
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                                                                                                                                                                                                                                                                                                                          disorder; Meckel's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA;
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Pred. No. 7.2e+
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                                                                                                                                                                                                                                                                                                                            diverticulum;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   inoma - includes a a promoter and a g
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07-JUN-2000; 28-JUN-2000; 30-JUN-2000; 07-JUL-2000; 07-JUL-2000; 11-JUL-2000; 11-JUL-2000; 14-JUL-2000;

-MAR-2000; 3-APR-2000; 3-MAY-2000;

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13-OCT-2000;
20-OCT-2000;
20-NOV-2000;
20-DEC-2000;
20-DE
                                                                         Polynucleotides encoding digestive system antigens, useful for diagnosing, treating, preventing and/or prognozing disorders of digestive system, particularly cancer and cancer metastases.
                                                                                                                                      WPI;
The present invention provides the protein and coding sequences number of human digestive system antigens. These can be used in
                                              Disclosure;
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2000US-0241221P.
2000US-0241786P.
2000US-0241786P.
2000US-0241809P.
2000US-0246474P.
2000US-0246474P.
2000US-0246474P.
2000US-0246477P.
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2000US-0246747P.
2000US-0246524P.
2000US-0246524P.
2000US-0246524P.
2000US-0246524P.
2000US-0246528P.
2000US-0246513P.
2000US-0246513P.
2000US-0249211P.
2000US-024921P.
2000US-0251989P.
2000US-0251989P.
2000US-0251997P.
2000US-0259678P.
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                                              986pp;
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2000US-0190076P.
2000US-0198123P.
2000US-0209467P.
2000US-02154886P.
2000US-02154880P.
2000US-02154880P.
2000US-0217496P.
2000US-0217496P.
2000US-0217496P.
2000US-0224518P.
2000US-0225213P.
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2000US-02293143P.
2000US-0231443P.
2000US-0231444P.
2000US-0231443P.
2000US-0233063P.
2000US-0233063P.
2000US-0233637P.
20

26-JUL-2000;
26-JUL-2000;
14-AUG-2000;
12-AUG-2000;
12-AUG-2000;
13-AUG-2000;
13-AUG-2000;
10-SEP-2000;
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RESULT 268
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ID AAS320
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12-MAR-2000
11-MAR-2000
11-MAR-2000
11-MAY-2000
07-JUN-2000
07-JUN-2000
07-JUL-2000
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11-JUL-2000
11-JUL-2000
14-AUG-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      diagnosis, treatment and prevention of digestive system disorders, including cancer, Meckel's diverticulum, bacterial or parasitic infections, appendicitis, Hirschaprung's disease, chronic colitis or ulcerative colitis. The present sequence is a genomic DNA fragment encoding a digestive system antigen of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17-JAN-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TTTTTTGTATTTTTAGTAGAGATGGGGTTC 797
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           standard;
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nilarity 93.3%;
Conservative
   2000US
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                                                          S-0205515P.
S-0214886P.
S-0214880P.
S-0215135P.
S-0216880P.
S-0217487P.
S-0217496P.
S-0218290P.
S-0218290P.
S-0224519P.
S-0224519P.
S-0225214P.
S-0225214P.
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S-0189874P.
S-0190076P.
S-0198123P.
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Pred. No. 7.3e
0; Mismatches
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No. 7.
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.3e+02;
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   14-AUG-2000
14-AUG-2000
14-AUG-2000
18-AUG-2000
22-AUG-2000
23-AUG-2000
23-AUG-2000
01-SEP-2000
01-SEP
2000US-02270394

2000US-0229343P

2000US-0229343P

2000US-0229343P

2000US-0229343P

2000US-023943P

2000US-0231243P

2000US-0231243P

2000US-0231244P

2000US-0231249P

2000US-0231249P

2000US-0232399P

2000US-023239P

2000US-023368P

2000US-0235836P

2000US-0235836P

2000US-023682P

2000US-0241785P

2000US-0241785P

2000US-024186P

2000US-024647P

2000US-024647P

2000US-024647P

2000US-024647P

2000US-024647P

2000US-024652P

2000US-024652P
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08-NOV-2000;
08-NOV-2000;
08-NOV-2000;
08-NOV-2000;
08-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
                                                                                                                                                                                                                                                                                                                                        17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
01-DEC-2000;
01-DEC-2000;
05-DEC-2000;
05-DEC-2000;
05-DEC-2000;
06-DEC-2000;
08-DEC-2000;
08-DEC-2000;
08-DEC-2000;
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17-NOV-2000;
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17-NOV-2000;
                                                                                                                                                                                                                         used in preventing, treating particularly cancer of the li
                                                                                                                                                                                                                                   Isolated nucleic acid molecule encoding a human liver related protein used in preventing, treating or ameliorating disorders of the liver
                                                                                                                                                                                                                                                             WPI; 2001-457728/49.
                                                                                                                                                                                                                                                                                               (HUMA-) HUMAN GENOME SCI INC
                                                                                                                                                                                                                                                                              Ç
                                                                                                                                                                                                                                                                              Barash SC,
                                                                                                                                                                                                                                                                                                              2000US-024652P

2000US-024661P

2000US-024661P

2000US-024661P

2000US-024961P

2000US-024921P

2000US-024924P

2000US-024924P

2000US-024924P

2000US-024924P

2000US-024924P

2000US-024924P

2000US-024924P

2000US-024928P

2000US-025198P

2000US-025198P

2000US-025199P

2000US-025199P
                                                                                                                                                                                                                            Liver.
                                                                                                                                                                                                                                                                                MS
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Claim

28

SEQ ID NO 551; 526pp; English.

CC humans, mice, rabbits, goats, horses, cats, dogs, chickens or sheep. A CC pathological condition can be determined by detecting the presence or absence of a mutation in a liver associated polynucleotide. The treatable disorders include autoimmune diseases such as rheumatoid arthritis, CC disorders include autoimmune diseases such as rheumatoid arthritis, CC cardiovascular disorders such as cardiac arrest, cerebrovascular CC disorders such as cardiac arrest, cerebrovascular CC disorders such as cerebral ischaemia, nervous system disorders such as CC disorders such as corneal infection, endocrine disorders such as CC coular disorders such as corneal infection, endocrine disorders such as CC cremature labour and infertility, gastrointestinal disorders such as CC crohn's disease, renal disorders such as glomerulonephritis and CC respiratory disorders such as asthma and pleurisy. The polypeptides can CC candination organs before transplantation, to regenerate tissues and in CC chemotaxis. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly Sequences AAS31827-AAS32182 represent genomic DNA molecules, which extends in the liver associated polypeptides of the invention. Liver associated polypeptides and their associated polynucleotides are useful in the diagnosis, treatment and prevention of various types of disorders in humans, mice, rabbits, goats, horses, cats, dogs, chickens or sheep. at ftp.wipo.int/pub/published_pct_sequences in e. ep. A encode

Sequence 32 ВÞ; 7 P W 0 .7 9 15 Η, 0 U; 0 Other;

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RESULT 269
ABN90430
ID ABN904
XX ABN904
XX ABN904
XX ABN904
XX Human
XX Homo s
YN US2000
XX Homo s
XX Homo s
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Best Local Similarity
Matches 28; Conserv
11-JUL-2000;

11-JUL-2000;

14-JUL-2000;

26-JUL-2000;

14-AUG-2000;

15-SEP-2000;

01-SEP-2000;

01-SEP-2000;

01-SEP-2000;

01-SEP-2000;

25-SEP-2000;

27-SEP-2000;

29-SEP-2000;

29-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; liver antigen; liver disorder; hepatic disorder; infection; hepatitis; viral; parasitic; bacterial; fungal; inflammatory condition; cirrhosis; granulomatous hepatitis; toxin damage; drug damage; autoimmune disease; wilson's disease; primary biliary cirrhosis; neoplastic disorder; cancer; tumour; portal hypertension; gastrointestinal disorder; hepatitis; drug screening; gene therapy; chromosome mapping; forensic analysis; antibody preparation; hepatotropic; cytostatic; antiinflammatory; virucide; antibacterial; hepatotropic; cytostatic; antiinflammatory; virucide; antibacterial;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABN90430;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human liver antigen HLDAV38 genomic sequence, SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         04-FEB-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17-JAN-2001; 2001US-00764887
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US2002042096-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      fungicide;
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  2000US-0216880P.
2000US-0217487P.
2000US-0217496P.
2000US-022963P.
2000US-02294519P.
2000US-0225268P.
2000US-0225267P.
2000US-022547P.
2000US-0225758P.
2000US-0225758P.
2000US-0225758P.
2000US-0225758P.
2000US-0225838P.
2000US-0229345P.
2000US-0229345P.
2000US-0229345P.
2000US-0229345P.
2000US-0239345P.
2000US-0239347P.
2000US-0239345P.
2000US-0234274P.
2000US-0234274P.
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2000US-0235838P.
2000US-0235838P.
2000US-0236368P.
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2000US-0214886P.
2000US-0216647P.
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Pred. No. 7.3e
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CC ABP40975) and to cDNAs encoding them (ABP90016-ABP90180), and also compasses polypeptides 90% identical and polynucleotides 95% identical compasses polypeptides 90% identical and polynucleotides 95% identical compasses polypeptides 90% identical and polynucleotides 95% identical compasses to the sequences of the invention. The invention additionally relates to creombinant vectors and host cells comprising human liver antigen composing or preventing various disorders in diagnosing, treating, compressing or preventing various disorders in diagnosing, treating, compassing or preventing various disorders in diagnosing, treating, compassing or preventing various disorders in diagnosing, treating, compassing or preventing various disorders in the spatitis C virus, bepatities C virus, bepatities C virus, conditions (e.g., Clonorchis sinensis, Echinococcus granulosus and compassing or toxins, c
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Best Local S
Matches 28
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08-DEC-2000;
08-DEC-2000;
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02-OCT-2000;
02-OCT-2000;
02-OCT-2000;
02-OCT-2000;
02-OCT-2000;
02-OCT-2000;
20-OCT-2000;
20-OCT-2000;
20-OCT-2000;
11-NOV-2000;
01-NOV-2000;
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(RUBE/)
(BARA/)
                                                                                                                                                Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New nucleic acid encoding human liver antigens, useful for diagnosis, treatment and prevention of e.g. hepatitis and hepatic cancer, also
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2002-381944/41.
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                                 768
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) RUBEN S M.
) BARASH S C.
                                                                        28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          nt and prevention of e.g. hep polypeptides and antibodies.
                                                                                        Similarity
                       TTTTTTGTATTTTTAGTAGAGATGGGGTTC 797
                                                                                                                                                  32
TTTTTGTATTTTAGTAGAGACAGGGTTC
                                                                                                                                                                                      USPTO
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2000US-0237037P.
2000US-0237033P.
2000US-0237039P.
2000US-0237049P.
2000US-0237049P.
2000US-0240960P.
2000US-0241809P.
2000US-0241809P.
2000US-0241809P.
2000US-0241809P.
2000US-0251869P.
                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                relates to 145 novel human liver antigens (ABP40831-to cDNAs encoding them (ABN90036-ABN90180), and also
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                                                                                                                                                P
                                                                                        2.7%;
93.3%;
                                                                                                                                                3 C; 7 G; 15
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                                                                      0;
                                                                      Score 26.8; DB 1
Pred. No. 7.3e+02
0; Mismatches
                                                                                                                                                T; 0
                                                                                                                                                  U; 0 Other;
30
                                                                                                           DB 1;
                                                                                                         Length 32;
                                                                        Indels
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                                                                        Gaps
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RESULT 270 ADJ15343 ID ADJ153

ADJ15343 standard; DNA; 32

BP

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14 AUG 2000

14 AUG 2000

14 AUG 2000

18 AUG 2000

22 AUG 2000

22 AUG 2000

23 AUG 2000

30 AUG 2000

01 SEP 2000
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14-AUG-2000;
14-AUG-2000;
14-AUG-2000;
14-AUG-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                liver; virucide; fungicide; antibacterial; antiparasitic; hepatotropic; antiinflammatory; cycostatic; litholytic; antiirheumatic; antiiarthritic; neuroprotective; antidiabetic; anticoagulant; thrombolytic; antiarteriosclerotic; cardiant; haemostatic; antiarrhythmic; ophthalmological; antiarteriosclerotic; vasotropic; osteopathic; ophthalmological; antiarteriosclerotic; vasotropic; osteopathic;
                                                                                                                                                                                                                                                                                                                                                        14-AUG-2000;
14-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                      14-AUG-2000;
14-AUG-2000;
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26-JUL-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 11-JUL-2000;
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2000US-0205515P

2000US-0214886P

2000US-0214886P

2000US-0215135P

2000US-02151880P

2000US-0217487P

2000US-0218290

2000US-0218290

2000US-0229541P

2000US-0225213P

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2000US-0225757P

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2000US-0189874P
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-SEP-2000;

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RESULT 271
AAQ77890/c
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                                                                                                                                                                                                                                                                                                                                                                             cc antiarrhythmic, ophthalmological, antiarreriosclerotic, vasotropic, cstopathic, nootropic, antiparkinsonian, anticonvulsant, neuroleptic, cstopathic, nootropic, antiparkinsonian, anticonvulsant, neuroleptic, cc vasotropic, cytostatic and gynaecological activities. The polypeptides cc and polynucleotides of the invention may be useful for diagnosis, controlled the invention of disorders of the liver such as controlled the invention of disorders of the liver such as controlled the invention of disorders of the liver such as controlled the invention of disorders, carriovascular disorders, pancreatic disorders, yallbladder consecutive disorders, immune disorders, blood related disorders, hyperproliferative clasorders, cardiovascular disorders, respiratory disorders, cardiovascular disorders, neurological diseases, endocrine consciuskeletal system disorders, neurological diseases, endocrine consciustive system disorders or developmental and inherited clisorders. The current sequence is that of the human liver-related consciusion of the invention. The current sequence is not shown within the specification per se but was obtained electronically from the USPTO
                                                                                                                                                                                                                                                                           Query Match 2.7%;
Best Local Similarity 93.3%;
Matches 28; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17-NOV-2000
17-NOV-2000
17-NOV-2000
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06-DEC-2000
08-DEC-2000
                                                                                                                    7890/c
AAQ77890
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a novel isolated, liver related polypeptide. The polypeptide of the invention demonstrates virucide, fungicide, antipacterial, antiparasitic, hepatotropic, antinflammatory, cyrostatic, litholytic, antirheumatic, antiarthritic, neuroprotective, antidiabetic, anticoagulant, thrombolytic, antiarteriosclerotic, cardiant, haemostatic, anticoagulant, thrombolytic, antiarteriosclerotic, cardiant, haemostatic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New liver related polypeptide, useful for diagnosis, treatment and/or prevention of liver, gastrointestinal, pancreatic, immune, blood related, endocrine, reproductive, hyperproliferative or reproductive disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2003-765398/72.
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                                  25-MAR-2003
06-JUL-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; SEQ ID NO 551; 181pp;
Neural thread protein AD10-7 cDNA 5' antisense oligonucleotide.
                                                                                     AAQ77890;
                                                                                                                                                                                                                                                                                                                                                                  the specification web-site.
                                                                                                                                                                                                                                            768
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                                                                                                                       standard; cDNA; 30
                                                                                                                                                                                                           TTTTTGTATTTTAGTAGAGACAGGGTTC 30
                                                                                                                                                                                                                             TTTTTGTATTTTTAGTAGAGATGGGGTTC 797
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2000US-0249265P.
2000US-024929P.
2000US-0250160P.
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2000US-0251989P.
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2000US-0251866P.
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2000US-0254097P.
2001US-0259678P.
2001US-00764887.
                                    (revised)
(first entry)
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                                                                                                                                                                                                                                                                              Score 26.8; D
Pred. No. 7.3e
0; Mismatches
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7.3e+02;
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2000US-0237039P.
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2000US-023703P.

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2000US-0231243P.
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2000US-0232399P.
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2000US-023427499P.
2000US-023427499P.

20-OCT-2000; 20-OCT-2000; 20-OCT-2000; 20-OCT-2000; 20-OCT-2000; 20-OCT-2000; 20-OCT-2000; 20-OCT-2000; 01-NOV-2000; 08-NOV-2000; 17-NOV-2000; 17-NOV-2000;

2000US-024050P.
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Detection of neural thread proteins - to detect sporadic and familial Alzheimer's disease, neuroectodermal tumours, malignant astrocytomas glioblastomas (Eng).
WPI; 1996-259865/26.
                                           De La Monte S,
                                                                                                                                                                                                                                                                                                                                               Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                binding fragment; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                              Neural thread protein; NTP; diagnosis; detection; Alzheimer's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Neural thread
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAT27744;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAT27744 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAQ77888-077890 are AD10-7 neural thread protein (NTP) antisense oligonucleotides, that can be used to down regulate or inhibit the expression of the NTP gene. These oligonucleotides could be used in the treatment of the folowing conditions Alzheimer's disease, neuroectodermal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 48; 158pp; English.
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                                                                                                                                                14-NOV-1994;
                                                                                                                                                                                              14-NOV-1995;
                                                                                                                                                                                                                                                  23-MAY-1996.
                                                                                                                                                                                                                                                                                              W09615272-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                            neuroectodermal
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                                                                                               GEN
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                                                                                               HOSPITAL CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP; 8 A; 4 C; 14 G; 4 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               protein antisense sequence
                                                                                                                                             94US-00340426
                                                                                                                                                                                                 95WO-US017111.
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                                              Wands JR;
                                                                                                                                                                                                                                                                                                                                                                                                                         tumour;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               astrocytomas and glioblastomas. (Updated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                       malignant
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Pred. No. 7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                         astrocytoma; monoclonal antibody;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 1;
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RESULT 273
AAH91474/c
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A method for detecting the presence of neural thread protein (NTP) having a molecular weight of 8, 14, 17, 21, 26 or 42 kD in a human subject comprises (a) contacting a sample from a human subject that is suspected of containing the NTP with at least one molecule capable of binding to the protein, and (b) detecting any of the molecule bound to the protein. The binding molecule is selected from an antibody free of natural impurities, a monoclonal antibody or a binding fragment of either of these. The method may be used for diagnosing the presence of Alzheimer's disease, neuroectodermal tumours and a malignant astrocytoma in a human. Expression of NTP nucleic acid may be inhibited using antisense oligonucleotides (See AAT27739-44)
                                                          Testing for the presence of polymorphisms bowel disease, using a hybridization assay
                                                                                                                                                                                                                                                                                                                                  misc_feature
                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                 chromosome
                                                                                                                                                                                                                                                                                                                                                                                                             Human; inflammatory bowel disease; Crohn's disease; ulcerative colitis; single nucleotide polymorphism; SNP; chromosome 19p13; paternity test;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             09-OCT-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Detection of neural thread protein in diagnosis also NTP DNA and protein sequences used in gene
                                                                                                  WPI; 2001-367874/38.
                                                                                                                          Daly M,
                                                                                                                                                   (WHED )
                                                                                                                                                                                        10-DEC-1999;
10-APR-2000;
                                                                                                                                                                                                                               11-DEC-2000; 2000WO-US033632
                                                                                                                                                                                                                                                                                  WC200142511-A2
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                                                                                                                                                     WHITEHEAD INST BIOMEDICAL ELLIPSIS BIOTHERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             standard; DNA;
                                                                                                                          Hudson TJ,
                                                                                                                                                                                                                                                                                                                                                                                                 5q31-33; forensic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                        99US-0170257P.
2000US-0196046P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
                                                                                                                                                                                                                                                                                                           note=
                                                                                                                                                                                                                                                                                                                       /*tag=
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                                                                                                                                                                                                                                                                                                                                                                                                            polymorphism;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2.7%;
                                                                                                                             Lander
                                                                                                                                                                                                                                                                                                          a
"SNP, optionally C
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                                                                                                                                                                                                                                                                                                                                                                                                 test; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 26.4;
Pred. No. 7
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                                                                                                                                                     CORP.
                                                                                                                             Rioux
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                                                                         associated with
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                                                                                                                             Siminovitch
                                                                                                                                                                                                                                                                                                                                                                                                                                                    polymorphic site
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                                                                                                                                                                                                                                                                                                           at this position"
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                                                                           inflammatory
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The present invention describes a method for polymorphisms associated with inflammatory

method for detecting the lammatory bowel diseases

the

e presence

of.

Claim 1; Page

62; 463pp; English.

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RESULT 274
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The invention provides polymorphic fragments of genes associated with hypertension. The nucleic acids including the polymorphic sites can be used as probes or primers for expressing variant proteins. Detection of the polymorphisms is useful in designing prophylactic and therapeutic regimes customized to underlying abnormalities. The polymorphisms can be used for association studies for hypertension, and in hypertension diagnostic assays. Where the polymorphisms have strong correlation with hypertension, within a gene, they are likely to have a causative role in hypertension. This information can be used to find the precise role of a polymorphism in the disease, and this can be used to identify potential drugs which combat the disease. The polymorphisms can be tested for association with other diseases e.g. agammaglobulinemia, diabetes insipidus, Lesch-Nyhan syndrome, muscular dystrophy, Wiskott-Aldrich syndrome, Fabrys disease, familial hypercholesterolemia, polycystic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                used
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Polymorphic fragment of hypertension associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAA04663 standard; DNA;
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03-MAY-1999;
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familial colonic polyposis; osteogenesis imperfecta; porphy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Lesch-Nyhan syndrome; muscular dystrophy; Wiskott-Aldrich syndrome;
Fabrys disease; familial hypercholesterolemia; hereditary spherocytosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Polymorphism; hypertension; agammaglobulinemia; diabetes insipidus;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   679 TGCAACCTCTGCCTCCCGGGTTCAAGTTATTC 710
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nilarity 87.5%;
Conservative
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99US-00304232.
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                                                                                                                                                                                                                                                                                                                                                                                         English.
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No. 7
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RESULT 275
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The invention provides polymorphic fragments of genes associated with CC hypertension. The nucleic acids including the polymorphic sites can be CC used as probes or primers for expressing variant proteins. Detection of CC the polymorphisms is useful in designing prophylactic and therapeutic CC regimes customized to underlying abnormalities. The polymorphisms can be CC diagnostic assays. Where the polymorphisms have strong correlation with CC hypertension, within a gene, they are likely to have a causative role in hypertension. This information can be used to find the precise role of a CC polymorphism in the disease, and this can be used to identify potential CC drugs which combat the disease, and this can be used to identify potential CC association with other diseases e.g. agammaglobulinemia, diabetes consipidus, Lesch-Nyhan syndrome, muscular dystrophy, Wiskott-Aldrich CC syndrome, Fabrys disease, familial hypercholesterolemia, polycystic kidney disease, hereditary spherocytosis, von Willebrands disease,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 29
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03-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Fabrys disease; familial hypercholesterolemia; hereditary spherocytosis; polycystic kidney disease; von Willebrands disease; forensic; human; tuberous sclerosis; hereditary hemorrhagica telangiectasia; familial colonic polyposis; osteogenesis imperfecta; porphyria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Polymorphism; hypertension; agammaglobulinemia; diabetes insipidus;
Lesch-Nyhan syndrome; muscular dystrophy; Wiskott-Aldrich syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAA03961 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Polymorphic fragment of hypertension associated gene APOC1.
                                                                                                                                                                                                                                                                                 Claim
                                                                                                                                                                                                                                                                                                                                   Novel nucleic acids containing polymorphisms used in the diagnosis
                                                                                                                                                                                                                                                                                                                                                                       WPI; 2000-107928/10
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UNIV CASE WESTERN RESERVE
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                                                                                                                                                                                                                                                                                 English.
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Pred. No.
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tuberous sclerosis, hereditary hemorrhagica telangiectasia, familial

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AAA03993
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         The invention provides polymorphic fragments of genes associated with the pretension. The nucleic acids including the polymorphic sites can be used as probes or primers for expressing variant proteins. Detection of the polymorphisms is useful in designing prophylactic and therapeutic regimes customized to underlying abnormalities. The polymorphisms can be used for association studies for hypertension, and in hypertension of diagnostic assays. Where the polymorphisms have strong correlation with hypertension, within a gene, they are likely to have a causative role in hypertension. This information can be used to find the precise role of a polymorphism in the disease, and this can be used to identify potential of drugs which combat the disease. The polymorphisms can be tested for association with other diseases e.g. agammaglobulinemia, diabetes can spociation with other diseases e.g. agammaglobulinemia, diabetes considered the process of t
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Lesch-Nyhan syndrome; muscular dystrophy; Wiskott-Aldrich
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UNIV CASE WESTERN RESERVE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Chakravarti A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               colonic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CCTCCCGGGTTCAASTGATTCTCCTGCC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 9 A; 5 C; 11 G; 3 T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                 acids containing polymorphisms used in the diagnosis of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       99US-00304232
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            98US-0084641P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           99EP-00250150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          polyposis; osteogenesis imperfecta;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA;
hereditary hemorrhagica telangiectasia,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2.6%;
92.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    of hypertension associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       29
                                                                                                                                                                                                                                                                                                                                                                                English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Haluska MK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ₽P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pred. No. 7.5
L; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 26;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  7.5e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           <u>بر</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        diabetes insipidus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          porphyria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APOC4.
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RESULT 278
AAT27743/c
ID AAT2774
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7743/c AAT27743

standard;

DNA;

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RESULT 277
AAQ77889/c
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Best Local S
                                                            Best
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                                               Matches
                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              colonic polyposis, Ehlers-Danlos syndrome, osteogenesis imperfecta, and acute intermittent porphyria. The polymorphic forms can also be used in forensics to identify individuals
                                                                                                                                             AAQ77888-Q77890 are AD10-7 neural thread protein (NTP) antisense oligonucleotides, that can be used to down regulate or inhibit the expression of the NTP gene. These oligonucleotides could be used in the treatment of the following conditions Alzheimer's disease, neuroectodermal
                                                                                                Sequence
                                                                                                                                                                                                         Disclosure; Page 48; 158pp; English.
                                                                                                                                                                                                                                   glioblastomas (Eng).
                                                                                                                                                                                                                                             Detection of neural thread proteins - to detect sporadic and familial Alzheimer's disease, neuroectodermal tumours, malignant astrocytomas
                                                                                                                                                                                                                                                                                    WPI; 1994-341497/42
                                                                                                                                                                                                                                                                                                             De La
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Neural thread protein AD10-7; Alzheimer's; neuroectodermal tumours; malignant astrocytomas; glioblastomas; 5' antisense therapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Neural thread
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  06-JUL-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             25-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAQ77889;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAQ77889
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 29 BP; 8 A; 5 C; 9 G; 6 T; 0 U; 1 Other;
                                                                                                                                  tumours,
                                                                                                                                                                                                                                                                                                                                                                                     20-APR-1994;
                                                                                                                                                                                                                                                                                                                                                                                                            27-OCT-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                     WO9423756-A1.
                                                                                                                                                                                                                                                                                                                                    (GEHO ) GEN HOSPITAL CORP
                                                            Local
                       980
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               860
                                                                                                                                                                                                                                                                                                             Monte SM,
29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        l Similarity
26; Conserv
                                               l Similarity 27; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 AAGTGCTAGGATTAYAGGCGTGAGCCAC 28
                                                                                                                      malignant astrocytomas and glioblastomas.
correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAGTGCTGGGATTACAGGCGTGAGCCAC 887
                        GCAACCTCTGCCTCCCGGGCTCAAGCGAT 1008
                                                                                                30 BP; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             standard; cDNA; 30
                                               2.6%;
nilarity 93.1%;
Conservative
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(first entry)
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                                                                                                A; 7 C; 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AD10-7
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Pred. No. 7.5e
1; Mismatches
                                                                                                G; 5 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cDNA 5' antisense oligonucleotide.
                                                           Score 25.8; DB 1;
Pred. No. 7.8e+02;
                                                Mismatches
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                                                                                                0 Other;
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                                                                      Length 30;
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                                                  Indels
                                                                                                                                    (Updated on 25-MAR-
                                               <u>,</u>
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Best Local S
Matches 27
                                                                                                                                                                                                                                                                                                                                                                                          A method for detecting the presence of neural thread protein (NTP) having a molecular weight of 8, 14, 17, 21, 26 or 42 kD in a human subject comprises (a) contacting a sample from a human subject that is suspected of containing the NTP with at least one molecule capable of binding to the protein; and (b) detecting any of the molecule bound to the protein. The binding molecule is selected from an antibody free of natural impurities, a monoclonal antibody or a binding fragment of either of these. The method may be used for diagnosing the presence of Alzheimer's disease, neuroectodermal tumours and a malignant astrocytoma in a human. Expression of NTP nucleic acid may be inhibited using antisense oligonucleotides (See ANT27739-44)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Neural thread protein; NTP; diagnosis; detection; Alzheimer's disease; neuroectodermal tumour; malignant astrocytoma; monoclonal antibody; binding fragment; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Neural
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14-NOV-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 48; 238pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Detection of neural thread protein in diagnosis of Alzheimer's also NTP DNA and protein sequences used in gene and anti:sense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1996-259865/26.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             14-NOV-1994;
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                                                                                                                                      25-MAR-2003
25-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO9615272-A1
                                                                       Enhancer element; carcinoma; tumor; cancer; SLPI gene; secretory leukoprotease-inhibitor gene; cytokeratin ge
                                                                                                               Enhancer element er-3 conserved basepair sequence.
                                                                                                                                                                                                    AAQ73573 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (GEHO ) GEN HOSPITAL CORP.
             misc_difference
                                                   Homo
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                                                                                                                                                                                                                                                                                            980
                                                  sapiens
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                                                                                                                                                                                                                                                                                                                  l Similarity
27; Conserv
                                                                                                                                                                                                                                                                                   GCAACCTCTGCCTCCCGGGCTCAAGCGAT 1008
                                                                                                                                                                                                                                                                                                                                                                      30
                                                                                                                                                                                                                                                                    GCAACCTCCGCCTCCCGGGTTCAAGCGAT 1
                                                                                                                                                                                                                                                                                                                                                                     ВP;
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                                                                                                                                                                                                                                                                                                                  2.6%;
ilarity 93.1%;
Conservative
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                                                                                                                                      (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             94US-00340426
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                                                                                                                                                                                                                                                                                                                                                                     5 A;
           Location/Qualifiers
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/*tag=
                                                                                                                                                                                                    DNA;
                                                                                                                                                                                                                                                                                                                                                                      7 C; 13 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                    31
                                                                                                                                                                                                      ВP
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                                                                                                                                                                                                                                                                                                                                Score 25.8;
Pred. No. 7
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                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                  .8e+02
                                                                                                                                                                                                                                                                                                                                              DB 1;
                                                                                                                                                                                                                                                                                                                                             Length 30;
                                                                                                                                                                                                                                                                                                                        Indels
                                                                          gene-8;
                                                                                                                                                                                                                                                                                                                     ٥,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              therapy.
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RESULT 280
AAA78748/c
ID AAA787
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Query Match
Best Local &
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This enhancer element is part of a DNA construct used for treating human carcinoma which contains a cancer therapeutic protein under the control of a promoter and 3 enhancer sequences in a specific 5'-3' order. This enhancer element is derived from the flanking region of the human epithelial cell secretory leukoprotease-inhibitor gene. (Updated on 25-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              misc_difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sequences.
                                                                                                                                                                                                                                                                                                                                                                               Human; genomic DNA; polymorphism; genome; allele-specific; primer; hybridisation; polymorphic site; forensic; paternity testing; medically testing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAA78748
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 31 BP; 7 A; 10 C; 7 G; 5 T; 0 U; 2 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; Fig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA construct for treating therapeutic gene under the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1994-316537/39.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Garver RI,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   24-MAR-1993;
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                                                                                                                                                                                                                                                                              EP1024200-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human genomic DNA polymorphic site sequence tag
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20-NOV-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAA78748;
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                    WPI; 2000-500198/45
                                                               Patil N,
                                                                                                                                                27-JAN-1999;
                                                                                                                                                                                        26-JAN-2000; 2000EP-00250023
                                                                                                                                                                                                                                   02-AUG-2000.
                                                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                               phenotypic trait; genetic
                                                                                                       (AFFY-)
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                                                                                                          AFFYMETRIX INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      standard;
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                                                               Shah N,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  6; 54pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
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/label=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA;
                                                               Warrigton
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oter and a gp. of enhancer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CC base, or the complement of the segment. Also described are: (1) an allele CC -specific oligonucleotide that hybridises to a segment of the novelty; (2) an isolated nucleic acid comprising a sequence of the novelty where CC the polymorphic site within the sequence is occupied by a base other than CC the reference base indicated in the specification; and (3) analysing a CC nucleic acid, comprising obtaining a nucleic acid from an individual, and CC determining a base occupying any one of the polymorphic sites of the CC novelty. The nucleic acid segments and method can be used to analyse an CC individuals nucleic acid segments and method can be used to analyse an CC method can also be used to test for a disease phenotype and correlate the presence of the phenotype with a particular polymorphism. The presence of polymorphisms with phenotypic traits and for genetic CC mapping of phenotypic traits. AAA78631 to AAA79262 represent sequence CC tags of human genomic DNA fragments containing polymorphic sites. The Dase occupying the polymorphic site is indicated using IUPAC-IUB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local :
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Generating five prime biased tandem tag libraries of cDNAs by isolating sample of mRNAs, amplifying the released tags, concatenating the amplified tags to form concatenated tags, amplifying and isolating the
                                                                                                                                                                                                                                                                                                                                                                                                                      06-MAR-2002; 2002US-00092885
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        09-OCT-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human tandem tag
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
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                                                                                                            WPI; 2003-831617/77.
                                                                                                                                                                                                                                                                                                                                                                   06-MAR-2002; 2002US-00092885
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                                                                                                                                                                                                            HERMIDA L C. HOPPA N L. JOHE K K.
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Matches
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            The invention provides polymorphic fragments of genes associated with hypertension. The nucleic acids including the polymorphic sites can be used as probes or primers for expressing variant proteins. Detection of the polymorphisms is useful in designing prophylactic and therapeutic regimes customized to underlying abnormalities. The polymorphisms can be used for association studies for hypertension, and in hypertension diagnostic assays. Where the polymorphisms have strong correlation with hypertension, within a gene, they are likely to have a causative role in hypertension. This information can be used to find the precise role of a polymorphism in the disease, and this can be used to identify potential
hypertension. This information opolymorphism in the disease, and drugs which combat the disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 mRNAs, amplifying the released tags, concater form concatenated tags, amplifying and isolat The present sequence is human tandem tag DNA
                                                                                                                                                                                                                                                                                                                                                                      07-MAY-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               polycystic kidney disease;
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tuberous sclerosis; hereditary hemorrhagica telangiectasia;
familial colonic polyposis; osteogenesis imperfecta; porphy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Fabrys disease; familial hypercholesterolemia; hereditary spherocytosis;
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                                                                                                                                                                                                                                                                                  JB,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               standard;
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                                                                                      The invention provides polymorphic fragments of genes associated with hypertension. The nucleic acids including the polymorphic sites can be used as probes or primers for expressing variant proteins. Detection of the polymorphisms is useful in designing prophylactic and therapeutic regimes customized to underlying abnormalities. The polymorphisms can be used for association studies for hypertension, and in hypertension diagnostic assays. Where the polymorphisms have strong correlation with hypertension, within a gene, they are likely to have a causative role in hypertension. This information can be used to find the precise role of a
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03-MAY-1999;
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familial colonic polyposis; osteogenesis imperfecta; porphyria;
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polymorphism in the disease, and this can be used to identify potential drugs which combat the disease. The polymorphisms can be tested for association with other diseases e.g. agammaglobulinemia, diabetes
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AFFYMETRIX INC.
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                                                                                                                                                                                                                                                                                                                                                                                                     Page 24;
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             hypertension. The nucleic acids including the polymorphic sites can be used as probes or primers for expressing variant proteins. Detection of the polymorphisms is useful in designing prophylactic and therapeutic regimes customized to underlying abnormalities. The polymorphisms can be used for association studies for hypertension, and in hypertension with diagnostic assays. Where the polymorphisms have strong correlation with hypertension, within a gene, they are likely to have a causative role in hypertension. This information can be used to find the precise role of a polymorphism in the disease, and this can be used to identify potential drugs which combat the disease. The polymorphisms can be tested for association with other diseases e.g. agammaglobulinemia, diabetes association with other diseases e.g. agammaglobulinemia, diabetes
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disease, familial hypercholesterolemia,
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The invention provides polymorphic fragments of genes associated with hypertension. The nucleic acids including the polymorphic sites can be used as probes or primers for expressing variant proteins. Detection of the polymorphisms is useful in designing prophylactic and therapeutic regimes customized to underlying abnormalities. The polymorphisms can be used for association studies for hypertension, and in hypertension diagnostic assays. Where the polymorphisms have strong correlation with hypertension, within a gene, they are likely to have a causative role in hypertension. This information can be used to find the precise role of a polymorphism in the disease, and this can be used to identify potential drugs which combat the disease. The polymorphisms can be tested for association with other disease. The polymorphisms can be tested for insiphidus, Lesch-Nyhan syndrome, muscular dystrophy, Wiskott-Aldrich syndrome, Fabrys disease, familial hypercholesterolemia, polycystic
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03-MAY-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1; Page 39; 53pp;
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Jesch-Nyhan syndrome; muscular dystrophy; Wiskott-Aldrich syndrome;
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The invention provides polymorphic fragments of genes associated with hypertension. The nucleic acids including the polymorphic sites can be used as probes or primers for expressing variant proteins. Detection of the polymorphisms is useful in designing prophylactic and therapeutic regimes customized to underlying abnormalities. The polymorphisms can be used for association studies for hypertension, and in hypertension with hypertension, within a gene, they are likely to have a causative role in hypertension. This information can be used to find the precise role of a polymorphism in the disease, and this can be used to identify potential curves which combat the disease. The polymorphisms can be tested for association with other disease e.g. agammaglobulinemia, diabetes in syndrome, Fabrys disease, familial hypercholesterolemia, polycystic kidney disease, hereditary spherocytosis, von Willebrands disease, familial hypercholesterolemia, familial
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(UYCA-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Lesch-Nyhan syndrome; muscular dystrophy; Wiskott-Aldrich syndrome; Fabrys disease; familial hypercholesterolemia; hereditary spherocytosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Polymorphic fragment of hypertension associated gene TBXA2R
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     07-MAY-1998;
03-MAY-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EP955382-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Polymorphism; hypertension; agammaglobulinemia; diabetes insipidus;
tuberous sclerosis, hereditary hemorrhagica telangiectasia, colonic polyposis, Ehlers-Danlos syndrome, osteogenesis imp
                                                                                                                                                                                                                                                                                                Claim 1; Page 43; 53pp;
                                                                                                                                                                                                                                                                                                                              nypertension.
                                                                                                                                                                                                                                                                                                                                            Novel nucleic acids containing polymorphisms used in the diagnosis of
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99US-00304232.
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                      SNP flanking sequence, the SNPE primer is used as a genotyping primer. The oligonucleotides are useful for genotyping a nucleic acid sample by performing a single-nucleotide primer extension reaction. The oligonucleotides are useful for determining the presence, absence or identity of a SNP and for genotyping nucleic acid samples, for e.g. to assess by association analysis the genotype of an individual or group of individuals, having a pathological phenotypic trait suspected of being caused by one or more SNPs. Phenotypic traits include diseases e.g. agammaglobulinaemia, diabetes insipidus, Lesch-Nyhan syndrome, muscular dystrophy, familial hypercholesterolaemia, polycystic kidney disease, osteogenesis imperfecta and acute intermittent porphyria. Phenotypic traits also include symptoms of or susceptibility to multifactorial disease of which a component is or may be genetic such as autoimmune disease, including, rheumatoid arthritis, multiple sclerosis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     includes kits for determining the presence or absence of a SNP, using the oligonuclectides of the invention. The PCR primers are used to amplify a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide primer extension (SNPE) primers, and the sequences of regions flanking sites of single nucleotide polymorphisms SNPs. The present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 1; Page 59; 83pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Picoult-Newburg L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO200129262-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis; inflammation; forensic investigation; paternity analysis; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Single nucleotide polymorphism; SNP; single nucleotide primer extension;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SNP specific upper PCR primer SEQ ID 1785.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAH38989
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15-OCT-1999;
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infection by pathogenic
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Sequence 30 BP; 6 A; 1 C; 7 G; 16 T; 0 U; 0 Other;
                                                          paternity analysis. The present sequence
                                                                          microorganism. The method is also useful
                                      for a human
                                      SNP containing
                                      DNA sequence
                                                          represents a PCR primer specific
                                                                            in forensic investigations and
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Matches Query Match Best Local 26; Similarity Conservative 2.6%; 0 Score 25.4; DB 1 Pred. No. 8.2e+02 Mismatches DB 1; Length Indels 0, Gaps 0

밁 Ś 767 TTTTTTTGTATTTTTAGTAGAGATGGG 4 TITTTTGTATTTTAGTAGAGACGGG 30 793

AAH40734 standard; DNA; 30

14-AUG-2001 (first entry)

SNP specific lower PCR primer SEQ ID

Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis; inflammation; forensic investigation; paternity analysis; PCR primer; ss

Homo sapiens.

WO200129262-A2

26-APR-2001.

L3-OCT-2000; 2000WO-US028436

15-OCT-1999;

99US-0160096P.

(ORCH-) ORCHID BIOSCIENCES INC

Picoult-Newburg L,

acid New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nuc sample.

Claim 1; Page 68; 83pp; English

SNP flanking sequence, the SNPE primer is used as a genotyping primer. The oligonucleotides are useful for genotyping a nucleic acid sample by performing a single-nucleotide primer extension reaction. The oligonucleotides are useful for determining the presence, absence or identity of a SNP and for genotyping nucleic acid samples, for e.g. to assess by association analysis the genotype of an individual or group individuals, having a pathological phenotypic trait suspected of being caused by one or more SNPs. Phenotypic traits include diseases e.g. agammaglobulinaemia, diabetes insipidus, Lesch-Nyhan syndrome, muscula: dystrophy, familial hypercholesterolaemia, polycystic kidney disease, osteogenesis imperfecta and acute intermittent porphyria. Phenotypic traits also include symptoms of or susceptibility to multifactorial disease of which a component is or may be genetic such as autoimmune afficered. includes kits for determining the presence or absence of a SNP, using the oligonucleotides of the invention. The PCR primers are used to amplify a primer extension (SNPE) primers, and the sequences of regions flanking sites of single nucleotide polymorphisms SNPs. The present invention Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide arthritis, muscular γď 엺

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                                                                                                                            Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 30
                                                                                                                                                                                                                                                                             Alu 1 and Alu 2 probes were used in hybridisations carried out in an aq. medium comprising a cocktail of: 10% chondroltin sulphate A; 45% medium comprising a cocktail of: 10% chondroltin sulphate A; 45% furnamide;5X saline citrate;25mM phosphate; & 250 micro-g/ml sheared herring sperm DNA. The probes were chemically labelled with 3-4 biotin molecules per probe at the 3'termini. Excellent staining of the DNA of human cell nuclei resulted when either of the Alu 1 or Alu 2 probes were present at 60 ng/ml (or each was present at 30 ng/ml) of the probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US5116727-A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Increasing hybridisation rate between complementary polynucleotide - using water-soluble hetero:polysaccharide with sulphated N-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         31-AUG-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            31-AUG-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hybridisation
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                                                                                                                                                                                                    Sequence 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           acetyl:galactosamine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Brigati DJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (INIZ-) INIZIATIVE MARITTIME 1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               26-MAY-1992.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Col 7; 6pp; English.
                                                   AGCCTCCCAAAGTGCTGGGATTACA 405
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                                                                                                                                                                                                       BP;
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                                                                                                   Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           89US-00404990
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                                                                                                                            2.5%;
                                                                                                                                                                                                    6 C; 7
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Pred. No.
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No. 8
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Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genotyping; agammaglobulinaemia; diabettes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis; inflammation; forensic investigation; paternity analysis; primer; ss.
                                                                                                                                                                                                                                                                                      New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SNP specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAH40799 standard; DNA; 25
                                                                                                                                                                                                                                                           Claim
                                                                                                                                                                                                                                                                              acid sample.
                                                                                                                                                                                                                                                                                                                    WPI; 2001-290930/30.
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                                                                                                                                                                                                                                                                                                                                                                                               13-OCT-2000; 2000WO-US028436.
                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200129262-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                            15-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                         (ORCH-) ORCHID BIOSCIENCES INC
                                                                                                                                                                                                                                                           1; Page 68; 83pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SNPE primer SEQ ID 3595
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
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                                                                                                                                                                                                                                                            English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP.
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Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide CC primer extension (SNPE) primers, and the sequences of regions flanking CC sites of single nucleotide polymorphisms SNPs. The present invention CC includes kits for determining the presence or absence of a SNP, using the CC oligonucleotides of the invention. The PCR primers are used to amplify a CC SNP flanking sequence, the SNPE primer is used as a genotyping primer. CC indigenucleotides are useful for genotyping a nucleic acid sample by CC performing a single-nucleotide primer extension reaction. The CC identity of a SNP and for genotyping nucleic acid sample by CC assess by association analysis the genotype of an individual or group of CC individuals, having a pathological phenotypic trait suspected of being CC caused by one or more SNPs. Phenotypic traits include diseases e.g. CC agammaglobulinaemia, diabetes insipidus, Leech-Nyhan syndrome, muscular CC osteogenesis imperfecta and acute intermittent porphyria. Phenotypic traits also include symptoms of or susceptibility to multifactorial CC diseases, including, rheumatoid arthritis, multiple scherosis, cancer, mervous system diseases and infection by pathogenic CC inflammation, cancer, nervous system diseases and infection by pathogenic CC inflammation. The method is also useful in forensic investigations and cCC enteroity analysis. The present sequence represents a single nucleotide primer extension (SNPE) primer specific for a human SNP containing DNA

Sequence 25 BP; 6 A; 4 C; 10 G; 5 T; 0 U; 0 Other;

맑 ş Matches Query Match Best Local (960 Similarity AAGTGCTGGGATTACAGGCGTGAGC 884 AAGTGCTGGGATTACAGGCGTGAGC 25 Conservative 2.5%; 0 Score 25; Pred. No. Mismatches DB 1; 7.6e+ 7.6e+02; 98 0; Length 25; Indels 0 Gaps 0

RESULT 292 AAH40799

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RESULT 293
AAS157
XX AAS157
XX AAS157
XX AAS157
XX 29-JAN
XX Human;
KW Human;
KW latent
KW latent
KW anti-r
XX 01-NOV
XX 01-NOV
XX 19-APR
XX (RERE-
XX Cloyd
XX Detect
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                                                                                                                                                                                                                                                                                                                                                                                                                              CC The invention relates to detecting integrated retroviruses in a human CC tissue sample comprising amplifying DNA in the sample with polymerase CC chain reaction (PCR) using one primer for an Alu sequence and the other CC primer for a retroviral sequence (e.g. human immunodeficiency virus 1 CC long terminal repeat, LTR), hybridising the PCR-amplified DNA with a CC probe that specifically recognises the amplified retroviral sequence and CC detecting hybridisation of the probe, where the hybridisation indicates CC detecting integrated retroviruses in human tissue sample such as paraffin CC embedded tissue sections or frozen tissue sections of lymphocytes, blood, CC rlymph nodes. The method quantitatively determines the number of cells with integrated HIV in the human tissue sample. The method is useful for CC determining and monitoring latent infection, preferably human CC immunodeficiency virus (HIV) latent infection in patients. The method conly detects integrated retroviruses, and thus allows accurate assessment CC of the frequency of productively and latently infected cells together in CC subjects undergoing anti-retroviruses, and the sample to the presence of latently infected cells so that they can conitor the presence of latently infected cells so that they can conitor the presence of latently infected cells so that they can conitor the presence of latently infected cells so that they can conitor the presence of latently infected cells so that they can conitor the presence of latently infected cells so that they can conitor the presence of latently infected cells so that they can conitor the presence of latently infected cells so that they can conitor the presence of latently infected cells so that they can conitor the presence of latently infected cells so that they can conitor the presence of latently infected cells so that they can conitor the presence of latently infected cells so that they can conitor the presence of latent sequence of latent sequence continued or not. The present conicon the contin
                                                                                                                                                                                Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Detecting integrated retroviruses in human sample, comprises amplifying DNA in sample using primers specific for Alu and retroviral sequences and detecting hybridization of probe recognizing amplified retroviral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-026164/03.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       29-JAN-2002
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                                                                                                                                                                                                                                                                                                                                                                                                               sequence is a human Alu sequence PCR primer used in the method
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                                                                                                                                                                                Similarity
                                                                         GCCTCCCAAAGTGCTGGGATTACAG 406
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                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                            BP; 6 A; 7 C; 7 G; 5 T; 0 U; 0 Other;
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RESULT 294

misc_feature

Location/Qualifiers

Homo sapiens.

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RESULT 295
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Determining the presence of neoplastic molecular markers, by identifying the presence of markers in host test sample using array of neoplastic molecular marker specific reagents and analyzing the array of the
                                                                                 Human; inflammatory bowel disease; Crohn's disease; ulcerative colitis; single nucleotide polymorphism; SNP; chromosome 19p13; paternity test;
                                                                                                                                                            09-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 25 BP; 7 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 1; Page 16; 41pp; English
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                                                                                                                             Human inflammatory bowel disease associated polymorphic site #673.
                                                                                                                                                                                                                    AAH91598 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (CEMI-) CEMINES LLC
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                                                                                                                                                                                                                                                                                            858 CAAAGTGCTGGGATTACAGGCGTGA 882
                                                                                                                                                                                                                                                                                                                                                                    Local
                                                                                                                                                                                                                                                                                                                                                  2.5%; Score 25;
1 Similarity 100.0%; Pred. No.
25; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PCR primer SEQ ID NO: 179
                                                                   polymorphism; SNP; chromosome 19p13; paternity test;
3; forensic test; gene therapy; ds.
                                                                                                                                                                                                                                                                                                                                                                                                               4 C; 9 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                       29
                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                      7.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                    DB 1;
                                                                                                                                                                                                                                                                                                                                                                                  Length 25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 296
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
Matches 26; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention describes a method for detecting the presence of polymorphisms associated with inflammatory bowel diseases such as ulcerative colitis and Crohn's disease. The methods can be used to detect the presence of genetic polymorphisms associated with inflammatory bowel disease and correlating their occurrence with disease states. They may be used in this way for phenotypic correlations, forensics, paternity testing, medicine and genetic analysis. The present sequence is a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Testing for the bowel disease,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10-DEC-1999; 99US-0170257P.
10-APR-2000; 2000US-0196046P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  14-JUN-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Polymorphism; hypertension; agammaglobulinemia; diabetes insipidus; lesch-Nyhan syndrome; muscular dystrophy; Wiskott-Aldrich syndrome; Fabrys disease; familial hypercholesterolemia; hereditary spherocytosis; polycystic kidney disease; von Willebrands disease; forensic; human; tuberous selerosis; hereditary hemorrhagica telangiectasia; tuberous selerosis; hereditary hemorrhagica telangiectasia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAA03985
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAA03985 standard; DNA; 29 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          polymorphic site described in the exemplification of the invention
                                              07-MAY-1998;
03-MAY-1999;
                                                                                                                                                                     07-MAY-1999;
                                                                                                                                                                                                                                                  10-NOV-1999.
                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ehlers-Danlos
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Polymorphic fragment of hypertension associated gene APOC3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      22-MAY-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      familial colonic polyposis; osteogenesis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (WHED )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                670
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TTGGCTCACTGCAACCTCTGCCTCCCGGG 698
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hudson TJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TTGGCTCACTGCAANCTCCACCTCCCGGG
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                                          98US-0084641P.
99US-00304232.
                                                                                                                                                                     99EP-00250150
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/note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2.5%;
89.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Lander ES,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          "SNP, optionally A or C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 24.8;
Pred. No. 8.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   .5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          imperfecta;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          porphyria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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RESULT 297
ABK65978
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention provides polymorphic fragments of genes associated with the phypertension. The nucleic acids including the polymorphic sites can be used as probes or primers for expressing variant proteins. Detection of the polymorphisms is useful in designing prophylactic and therapeutic regimes customized to underlying abnormalities. The polymorphisms can be used for association studies for hypertension, and in hypertension with diagnostic assays. Where the polymorphisms have strong correlation with hypertension, within a gene, they are likely to have a causative role in hypertension. This information can be used to find the precise role of a polymorphism in the disease, and this can be used to identify potential drugs which combat the disease. The polymorphisms can be tested for association with other diseases e.g. agammaglobulinemia, diabetes insipidus, Lesch-Nyhan syndrome, mascular dystrophy, wiskott-Aldrich cyndrome, Fabrys disease, familial hypercholesterolemia, polycystic chalcular disease, hereditary spherocytosis, von Willebrands disease, tuberous sclerosis, hereditary spherocytosis, von Willebrands disease, familial colonic polyposis, Bheers-Danlos syndrome, osteogenessis imperfecta, and acute intermittent porphyria. The polymorphic forms can also be used in forensics to identify individuals
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local S
Matches 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 29 BP; 6 A; 8 C; 10 G; 4 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1; Page 22; 53pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (UYCA-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           nypertension.
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(UYCA-) UNIV CASE WESTERN RESERVE.
                                                                                                                                                                05-JAN-1999;
                                                                                                                                                                                                                         US6352829-B1
                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                   Primer; ss;
                                                                                                                                                                                                                                                                                                              Human gene
                                                                                                                                                                                                                                                                                                                                                02-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                              ABK65978;
                                                                                                                                                                                                                                                                                                                                                                                                         ABK65978 standard;
                                                                        Chenchik A,
                                                                                                                                  21-MAY-1997;
                                                                                                                                                                                            05-MAR-2002.
                                                                                                      (CLON-) CLONTECH LAB
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                                                                                                                                                                                                                                                                                                               specific PCR primer #66.
                                                                                                                                                                                                                                                                                    DNA
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                                                                                                                                                                99US-00225928
                                                                                                                                    97US-00859998
                                                                                                                                                                                                                                                                                    microarray; differential expression analysis; human.
                                                                                                                                                                                                                                                                                                                                                                                                            DNA;
                                                                                                                                                                                                                                                                                                                                                entry)
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                                                                                                                                                                                                                                                                                                                                                                                                            26
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                                                                           Bibilashvilli
                                                                                                                                                                                                                                                                                                                                                                                                            ΒP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 24.6;
Pred. No. 8.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 29;
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Producing sub-population of differences in RNA profiles

labeled between

nucleic acids, useful for analyzing several different physiological

sources, using

set of distinct gene specific primers.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 298
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cc acids (NAS) comprising contacting a NA sample from a physiological course, with a pool of 50 distinct gene specific primers under suitable conditions to enzymatically generate sub-population of NAS, where each ceach labeled NA is generated using a single gene specific primer. The ceach labeled NA is generated using a single gene specific primer. The ceach labeled NA is generated using a single gene specific primer. The complete for analysing the differences in the RNA profiles between several control of the population of labeled NAS which is comprising the populations for each physiological sources, comprising the populations for each physiological sources, comprising the population, where the comparison is preferably confirmed by hybridising the labeled NAS for each of the distinct comparison of a substrate to produce a hybridisation pattern for each of the sources, and comparing the patterns for each of the sources, where confirmed is a confirmed by the patterns for each of the sources, where confirmed is a profile of diseased a normal tissue e.g. neoplastic a normal confirmed gene specific PCR primer used in the method of the printed confirmed confirmed this patent did not form part of the printed confirmation, but was obtained in electronic format directly from USPTO are betterned to the printed confirmed the confirmed by the patterned to the printed confirmed the confirmed by the printed confirmed the confirmed by the printed confirmed the confirmed by the printed confirmed by the printed
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Best Local
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                  Example 3; SEQ ID NO 1072; 11pp; English
                                                        producing sub-population of labeled nucleic acids, useful for analyzing differences in RNA profiles between several different physiological sources, using set of distinct gene specific primers.
                                                                                                                                                      WPI; 2002-314699/35.
                                                                                                                                                                                                Chenchik A,
                                                                                                                                                                                                                                                                                                                                     05-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                  05-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                              US6352829-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Primer; ss; DNA microarray; differential expression analysis; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human gene specific PCR primer #1072.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 02-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABK66984 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             at http.wipo.seqdata.uspto.gov/sequence.html?DocID=6352829B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 3; SEQ ID NO 66; 11pp; English.
                                                                                                                                                                                                                                            (CLON-) CLONTECH LAB INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           invention relates to producing a sub-population of labeled nucleic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 26 BP; 8 A; 4 C; 9 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAAGTGCTAGGATTACAGGCGTGAGC
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                                                                                                                                                                                                Jokhadze G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                           97US-00859998
                                                                                                                                                                                                                                                                                                                                        9908-00225928
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2.5%;
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                                                                                                                                                                                                  Bibilashvilli R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ₽₽
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Pred. No. 8.
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8.3e+02;
1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 299
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          different physiological sources, where the method comprises producing subpopulation of labeled NAs for the different physiological sources, comprising the populations for each physiological source to identify differences in the population, where the comparison is preferably performed by hybridising the labeled NAs for each of the distinct physiological sources to an array of probe NAs stably associated with the surface of a substrate to produce a hybridisation pattern for each of the sources, and comparing the patterns for each of the sources, where differential gene expression assays are utilised in differential expression analysis of diseased a normal tissue e.g. neoplastic a normal tissue, or different tissue or subtissue types. The present sequence is a human gene specific PCR primer used in the method of the invention. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from USPTO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         acids (NAs) comprising contacting a NA sample from a physiological source, with a pool of 50 distinct gene specific primers under suitable conditions to enzymatically generate sub-population of NAs, where each gene specific primer has a sequence complementary to a distinct mRNA, and each labeled NA is generated using a single gene specific primer. The method is useful for producing a sub-population of labeled NAs which is useful for producing a sub-population of labeled NAs which is useful for analysing the differences in the RNA profiles between several
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; inflammatory bowel disease; Crohn's disease; ulcerative colitis; single nucleotide polymorphism; SNP; chromosome 19p13: paternity test.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   09-OCT-2001 (first entry)
                                                                    Daly M,
                                                                                                                                                                                                                                                                                                                                                                        misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      chromosome 5q31-33; forensic test; gene therapy; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human inflammatory bowel disease associated polymorphic site #605.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAH91530;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAH91530 standard; DNA; 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          at http.wipo.seqdata.uspto.gov/sequence.html?DocID=6352829B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to producing a sub-population of labeled nucleic
                                                                                                         (WHED )
                                                                                                                                                               10-DEC-1999;
10-APR-2000;
                                                                                                                                                                                                                        11-DEC-2000; 2000WO-US033632
                                                                                                                                                                                                                                                             14-JUN-2001.
                                                                                                                                                                                                                                                                                                  WO200142511-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                650 TGGAGTGCAGTGGCGCAATCTTGGCT 675
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25;
                                                                                                         WHITEHEAD INST BIOMEDICAL RES. ELLIPSIS BIOTHERAPEUTICS CORP.
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                                                                    Hudson TJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TGGAGTGCAATGGCGCAATCTTGGCT 26
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nilarity 96.2%;
Conservative
                                                                                                                                                               99US-0170257P.
2000US-0196046P.
                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                     note= "SNP, optionally C or G at this position"
                                                                      Lander ES,
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Pred. No. 8
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                                                                      Rioux J,
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                                                                        Siminovitch K;
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Testing for the presence of polymorphisms associated with inflammatory

WPI; 2001-367874/38

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RESULT 300
AAA04000/c
ID AAA040
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Best Local S
Matches 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  polymorphisms associated with inflammatory bowel diseases such as ulcerative colitis and Crohn's disease. The methods can be used to detect the presence of genetic golymorphisms associated with inflammatory bowel disease and correlating their occurrence with disease states. They may be used in this way for phenotypic correlations, forensics, paternity testing, medicine and genetic analysis. The present sequence is a polymorphic site described in the exemplification of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  polycystic kidney disease; von Willebrands disease; forensi
tuberous sclerosis; hereditary hemorrhagica telangiectasia;
familial colonic polyposis; osteogenesis imperfecta; porphy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Polymorphism; hypertension; agammaglobulinemia; diabetes insipidus;
Lesch-Nyhan syndrome; muscular dystrophy; Wiskott-Aldrich syndrome;
Fabrys disease; familial hypercholesterolemia; hereditary spherocytosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 28 BP; 4 A; 9 C; 4 G; 10 T; 0 U; 1 Other;
The invention provides polymorphic fragments of genes associated with hypertension. The nucleic acids including the polymorphic sites can be used as probes or primers for expressing variant proteins. Detection of the polymorphisms is useful in designing prophylactic and therapeutic regimes customized to underlying abnormalities. The polymorphisms can be used for association studies for hypertension, and in hypertension diagnostic assays. Where the polymorphisms have strong correlation with hypertension, within a gene, they are likely to have a causative role in
                                                                                                                                                                                                                                                                                                                                 07-MAY-1998;
03-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22-MAY-2000
                                                                                                                                                                                      Novel nucleic acids containing polymorphisms used in the diagnosis
                                                                                                                                                                                                                       WPI; 2000-107928/10
                                                                                                                                                                                                                                                                                   (AFFY-)
(UYCA-)
                                                                                                                                                                                                                                                                                                                                                                                07-MAY-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ehlers-Danlos
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAA04000 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           present invention describes
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                                                                                                                                                                                                                                                                                     AFFYMETRIX INCUNIV CASE WEST
                                                                                                                                            Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TCAAGTGATTCTNCTGTCTCAGCCTCC 28
                                                                                                                                                                                                                                                      Chakravarti A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 fragment of hypertension associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           64; 463pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         using a hybridization
                                                                                                                                            22; 53pp;
                                                                                                                                                                                                                                                                                                                                 98US-0084641P
99US-00304232
                                                                                                                                                                                                                                                                                                                                                                                99EP-00250150
                                                                                                                                                                                                                                                                                     WESTERN RESERVE
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92.6%;
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                                                                                                                                            English.
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                                                                                                                                                                                                                                                      Haluska MK;
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Pred. No. 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       8.7e+02
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      forensic; human;
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07-MAY-1998; 07-MAY-1999;

98US-0084641P. 99US-00304232. 99EP-00250150 10-NOV-1999.

03-MAY-1999;

AFFYMETRIX INC. UNIV CASE WESTERN RESERVE.

The invention provides polymorphic fragments of genes associated with hypertension. The nucleic acids including the polymorphic sites can be used as probes or primers for expressing variant proteins. Detection of the polymorphisms is useful in designing prophylactic and therapeutic regimes customized to underlying abnormalities. The polymorphisms can be used for association studies for hypertension, and in hypertension diagnostic assays. Where the polymorphisms have strong correlation with hypertension, within a gene, they are likely to have a causative role in hypertension. This information can be used to find the precise role of a

Claim 1; Page 38; 53pp;

English.

Novel nucleic acids containing polymorphisms used in

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diagnosis

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WPI; 2000-107928/10

Fan (AFFY-) (UYCA-)

JB,

Chakravarti

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          hypertension. This information can be used to find the precise role of polymorphism in the disease, and this can be used to identify potential drugs which combat the disease. The polymorphisms can be tested for association with other diseases e.g. agammaglobulinemia, diabbtes insipidus, Lesch-Nyhan syndrome, muscular dystrophy, Wiskott-Aldrich syndrome, Fabrys disease, familial hypercholesterolemia, polycystic kidney disease, hereditary spherocytosis, von Willebrands disease, tuberous sclerosis, hereditary hemorrhagica telangiectasia, familial colonic polyposis, Ehers-Danlos syndrome, osteogenesis imperfecta, and acute intermittent porphyria. The polymorphic forms can also be used in forensics to identify individuals
                                                                                                                                                                                                                                                                                                                                                                                                                          Lesch-Nyhan syndrome; muscular dystrophy; Wiskott-Aldrich Fabrys disease; familial hypercholesterolemia; hereditary polycystic kidney disease; von Willebrands disease; forens
                                                                                                                                                                                                                                                                                                                                                                                                     polycystic kidney disease; von Willebrands disease; forensic; human; tuberous sclerosis; hereditary hemorrhagica telangiectasia; familial colonic polyposis; osteogenesis imperfecta; porphyria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAA04507 standard;
                                                                                                                                                                                                                                                                                                                                                EP955382-A2.
                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                               Ehlers-Danlos
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Polymorphism; hypertension; agammaglobulinemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Polymorphic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         fragment
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          5 C; 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          of hypertension associated
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Pred.
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No. 8.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      .9e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      diabetes
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Matches
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                     The invention provides polymorphic fragments of genes associated with hypertension. The nucleic acids including the polymorphic sites can be used as probes or primers for expressing variant proteins. Detection of the polymorphisms is useful in designing prophylactic and therapeutic regimes customized to underlying abnormalities. The polymorphisms can be used for association studies for hypertension, and in hypertension diagnostic assays. Where the polymorphisms have strong correlation with hypertension, within a gene, they are likely to have a causative role in hypertension. This information can be used to find the precise role of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-MAY-1998;
03-MAY-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Polymorphism; hypertension; agammaglobulinemia; diabetes insipidus;
Lesch-Nyhan syndrome; muscular dystrophy; Wiskott-Aldrich syndrome;
Fabrys disease; familial hypercholesterolemia; hereditary spherocytosis;
polycystic kidney disease; von Willebrands disease; forensic; human;
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                                                                                                                                                                                                                                                                                                                             Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EP955382-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                   Novel nucleic acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            tuberous sclerosis; hereditary hemorrhagica telangiectasia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (UYCA-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
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UNIV CASE WESTERN RESERVE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 colonic polyposis; osteogenesis anlos syndrome; ss.
                                                                                                                                                                                                                                                                                                                          Page 34; 53pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GGGATTACAGGTGTRAGCCACCGCGCCC 28
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
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99US-00304232.
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                                                                                                                                                                                                                                                                                                                                                                                                                containing polymorphisms used
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              29
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and this
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Pred. No. 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                imperfecta; porphyria;
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                                          drugs which combat the disease. The polymorphisms can be tested for association with other diseases e.g. agammaglobulinemia, diabetes insipidus, Lesch-Nyhan syndrome, muscular dystrophy, Wiskott-Aldrich syndrome, Fabrys disease, familial hypercholesterolemia, polycystic kidney disease, hereditary spherocytosis, von Willebrands disease, tuberous sclerosis, hereditary hemorrhagica telangictasia, familial colonic polyposis, Ehlers-Danlos syndrome, osteogenesis imperfecta, a acute intermittent porphyria. The polymorphic forms can also be used forensics to identify individuals
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Sequence
29 BP; 7 A; 9 C; 6 G; 6 T; 0 U; 1 Other;
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Score 24.4; DB 1
Pred. No. 8.9e+02
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                   AGCTGGGATTACGGGCACCTGCCACCAC
 AGCTGGGATTACASGCACCTGCCATCAC
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22-MAY-2000
                                    AAA03994 standard;
(first entry)
                                    DNA;
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Polymorphism; hypertension; agammaglobulinemia; diabetes insipidus; lesch-Nýhan syndrome; muscular dystrophy; Wiskott-Aldrich syndrome; Fabrys disease; familial hypercholesterolemia; hereditary spherocytpolycystic kidney disease; ron Willebrands disease; forensic; human Ehlers-Danlos tuberous sclerosis; hereditary hemorrhagica telangiectasia; familial colonic polyposis; osteogenesis imperfecta; porphyria; spherocytosis;

Polymorphic fragment of hypertension associated gene APOC4.

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EP955382-A2
                        Homo sapiens.
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ARASSILF 303
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03-MAY-1999;
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99US-00304232.
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Novel nucleic acids containing polymorphisms used in the diagnosis of.

WPI; 2000-107928/10

Chakravarti A,

Haluska MK

Claim 1; Page 22; 53pp; English

The invention provides polymorphic fragments of genes associated with hypertension. The nucleic acids including the polymorphic sites can be used as probes or primers for expressing variant proteins. Detection of the polymorphisms is useful in designing prophylactic and therapeutic regimes customized to underlying abnormalities. The polymorphisms can be used for association studies for hypertension, and in hypertension with diagnostic assays. Where the polymorphisms have strong correlation with hypertension, within a gene, they are likely to have a causative role in hypertension. This information can be used to find the precise role of a polymorphism in the disease, and this can be used to identify potential which combat the disease. The polymorphisms

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be used to

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Matches 25
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The invention provides polymorphic fragments of genes associated with hypertension. The nucleic acids including the polymorphic sites can be used as probes or primers for expressing variant proteins. Detection of the polymorphisms is useful in designing prophylactic and therapeutic regimes customized to underlying abnormalities. The polymorphisms can be used for association studies for hypertension, and in hypertension diagnostic assays. Where the polymorphisms have strong correlation with hypertension, within a gene, they are likely to have a causative role in hypertension. This information can be used to identify potential drugs which combat the disease, and this can be used to identify potential drugs which combat the disease. The polymorphisms can be tested for association with other diseases e.g. agammaglobulinemia, diabetes
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tuberous sclerosis; hereditary hemorrhagica telangiectasia;
familial colonic polyposis; osteogenesis imperfecta; porphyria;
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03-MAY-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2000-107928/10
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99US-00304232.
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                                                                                                                                                                                                                                                                                                                                                                                                                    English.
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Pred. No. 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               diagnosis
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RESULT 305
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         The invention provides polymorphic fragments of genes associated with hypertension. The nucleic acids including the polymorphic sites can be used as probes or primers for expressing variant proteins. Detection of the polymorphisms is useful in designing prophylactic and therapeutic regimes customized to underlying abnormalities. The polymorphisms can be used for association studies for hypertension, and in hypertension diagnostic assays. Where the polymorphisms have strong correlation with hypertension, within a gene, they are likely to have a causative role in hypertension. This information can be used to identify potential drugs which combat the disease. The polymorphisms can be tested for association with other diseases e.g. agammaglobulinemia, diabetes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Polymorphism; hypertension; agammaglobulinemia; diabetes insipidus; Lesch-Nyhan syndrome; muscular dystrophy; Wiskott-Aldrich syndrome; Fabrys disease; familial hypercholesterolemia; hereditary spherocytosis; polycystic kidney disease; von Willebrands disease; forensic; human; tuberous sclerosis; hereditary hemorrhagica telangiectasia;
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03-MAY-1999;
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                                                                                                                                                                                                               Claim 1; Page 32; 53pp; English.
                                                                                                                                                                                                                                                            Novel nucleic acids containing polymorphisms used in
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UNIV CASE WESTERN RESERVE.
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99US-00304232.
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                                                      (PSES) promoter construct, which contains the androgen receptor core region (AREC3) promoter of the prostate-specific membrane antigen (PSMA) enhancer (PSMEdel2) promoter of the PSMA gene. The PSES promoter construct of the invention is useful for reducing angiogenesis in prostate carcinoma cells and in targeting prostate carcinoma cells for destruction. The PSES promoter construct is also useful for identifying an agent that modulates PSES promoter activity. The present DNA sequence represents a PCR primer that was used in an example of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                    Novel chimeric PSES promoter construct comprising androgen receptor enhancer core region promoter of prostate-specific-antigen gene and PSME(del2) promoter of PSMA gene, useful for treating prostate cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              prostate-specific-antigen; PSA;
prostate-specific membrane antigen enhancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   chimeric; prostate-specific-enhancing sequence promoter;
androgen receptor core region promoter; AREC3 promoter;
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                                                                                                                                                                                                                                                                                     The invention comprises a chimeric prostate-specific-enhancing sequence (PSES) promoter construct, which contains the androgen receptor core
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LEE S.
KIM H.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Lee S,
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                                                                                                                                                                                                                                      500-1000bp fragments from an AluI-digest of human genomic DNA were ligated to M13mp19 RF DNA. E.coli JM109 were transformed by the ligation mixture. Filter replicates of the transformant colonies were screened with probe #1 and a second probe. The probes were derived from the Alu family consensus sequence. Phage which hybridised to both probes were plated at lower density and rescreened with the same probes. Singlestranded template DNA was extd. from cultures of these phage to isolate low-frequency repeat sequence probes LF1, LF2, LF3, LF19, LF20 and LF21. See also AAQ2913-Q29017 and AAQ29021-Q29038. (Updated on 25-MAR-2003 to
                                                                                                                                                                                                        Sequence 25 BP; 5 A; 8 C; 5 G; 5 T; 0 U; 2 Other;
                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 8; 41pp;
                                                                                                                                                                                                                                                                                                                                                                           New nucleic acid probes - have a labelled low frequency repetitive sequence for detecting overlaps among cloned DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EP505605-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Alu family consensus sequence-derived probe #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25-MAR-2003
23-FEB-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAQ29012;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAQ29012
                                                        AAH91549
                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1992-324992/40.
                                                                                                                                                                                                                                                                                                                                                                                                                                     Duncan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       11-APR-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30-SEP-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Low frequency
                                                                                                                                                                                                                                                                                                                                                                                                                                                           (UYWA-) UNIV WAYNE STATE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local
                                                                                                                                     863
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               673 GCTCACTGCAACCTCTGCCTCCGGG 698
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                                                                                                                                                                       Similarity
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                                                                                                                                                                                                                              PN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  standard;
                                                        standard;
                                                                                                                                     TGCTGGGATTACAGGCGTGAGCCAC 887
                                                                                                                 TGCTGGGATTACAGGYRTGAGCCAC
                                                                                                                                                                                                                              field.
                                                                                                                                                                                                                                                                                                                                                                                                                                     Solus JF,
                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                repeat; AluI restriction digest;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  91US-00676292
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       91EP-00105802
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA;
                                                        DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2.5%;
                                                                                                                                                                      2.4%;
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                                                                                                                                                                                                                                                                                                                                                       English.
                                                         ВP
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                                                                                                                                                           اد
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Pred. No. 9
                                                                                                                                                                        Pred. No.
                                                                                                                                                                                  Score 24.2;
                                                                                                                                                           Mismatches
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                                                                                                                                                                      8.2e+02
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                                                                                                                                                                                  DB 1;
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                                                                                                                                                                                Length 25;
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                                                                                                                                                            Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                88
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                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                        ligation
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Human

inflammatory bowel disease associated polymorphic site

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                                                                                                                                                                                   AAF29776
                                                                                                                                                                                               RESULT 309
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Best Local S
Matches 26
                                                                                                                                                                                                                                                                                                                                                                       The present invention describes a method for detecting the presence of polymorphisms associated with inflammatory bowel diseases such as ulcerative colitis and Conohn's disease. The methods can be used to detect the presence of genetic polymorphisms associated with inflammatory bowel disease and correlating their occurrence with disease states. They may be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human;
                                                                                                                                                                                                                                                                                                                                        used in this way for phenotypic correlations, forensics, paternity testing, medicine and genetic analysis. The present sequence is a polymorphic site described in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; Page 65; 463pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Testing for the bowel disease,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Daly M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10-DEC-1999; 99US-0170257P
10-APR-2000; 2000US-0196046P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO200142511-A2
                                                                                                                                                                         AAF29776 standard;
                                                                                                                                                                                                                                                                                                                   Sequence 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (ELLI-)
                                                                      preseniline-1;
                                                                                                          Preseniline-1
                                                                                                                               09-APR-2001
                                 WO200079000-A1
                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2001-367874/38
                                                                                                                                                                                                                                                    832
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     nucleotide polymorphism; SNP; chi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            inflammatory
                                                                                                                                                                                                                                μ.
                                                                                    PSEN1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WHITEHEAD INST BIOMEDICAL RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                             for the presence of polymorphisms associated with inflammatory isease, using a hybridization assay.
                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hudson TJ,
                                                                                                                                                                                                                                CATGTGATCTGCCNGCCTCAGCCTTCCAAA 30
                                                                                                                                                                                                                                                   CTTGTGATCTGCCTGCCTCGGCCTCCCAAA 861
                                                                                                                                                                                                                                                                                                                  BP; 6
                                                                                                                                                                                                                                                                         Conservative
                                                                                                                               (first entry)
                                                                          Alzheimer's chromosome
                                                                                                         gene promoter PCR primer
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/note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                    A;
                                                                                                                                                                          DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            bowel
                                                                                                                                                                                                                                                                                   2.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Lander ES,
                                                                                                                                                                                                                                                                                                                    11 C; 5 G; 7 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           a
"SNP,
                                                                                                                                                                          30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            disease; Crohn's disease; ulcerative colitis;
                                                                         disease; polymorphism;
14; PCR primer; ss.
                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            optionally T
                                                                                                                                                                                                                                                                                   Score 24.2; DB 1;
Pred. No. 9.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SNP; chromosome 19p13; paternity test; gene therapy; ds.
                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Rioux J,
                                                                                                           Prom6F.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             at this position"
                                                                                                                                                                                                                                                                                               Length 30
                                                                                     diagnosis;
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RESULT 310
AAH45830/c
ID AAH458
XX
AC AAH458
XX
      Query Match
Best Local S
Matches 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Determining wether a human subject has or is at -onset) Alzheimer's disease comprises detecting genetic lesion in the presenilin-1 gene.
The present invention describes a method for determining the length of telomeres, involving hybridising a chromosomal DNA extracted from a sample and a labeled DNA probe with a sequence complementary to a repetitive telomeric sequence, and measuring the labeled signal of the hybridised probe to give the size of telomere. This can be used to investigate tissue aging and the progression of cancer and in monitorir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention describes a method for determining the presence or susceptibility to Alzheimer's disease in humans, involving detecting genetic lesion in the presentiline-1 (PSENI) gene, found on chromosome 1 The genetic lesion is a polymorphism in the promoter or upstream regulatory region of the gene. The invention also describes transgenic animals which can be used to identify compounds useful in treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 1; Page 45; 56pp; English.
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                                                                                                                                                   Determining telomere size useful progression of cancer.
                                                                                                                                                                                                                                                                                                                                                                      JP2001095586-A.
                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Telomere size determination; chromosomal DNA; probe; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    11-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAH45830;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAH45830 standard; DNA; 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Alzheimer's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (VLAA-) VLAAMS INTERUNIVERSITAIR INST BIOTECHNOG
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                                                                                                                                                                                                     WPI; 2001-360495/38.
                                                                                                                                                                                                                                                                    30-SEP-1999;
                                                                                                                                                                                                                                                                                                    30-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                   repetitive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Telomere size determination
                                                                                                                    Example 2; Page 6; 8pp; Japanese.
                                                                                                                                                                                                                                   (IDET/)
                                                                                                                                                                                                                                                                                                                                     10-APR-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          205 GTCAGGCTGGTCTCGAACTCCCGACCTCA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 h 2.4%;
Similarity 89.7%;
26; Conservative
                                                                                                                                                                                                                                     IDE T.
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                                                                                                                                                                                                                                                                                                                                                                                                                                     sequence; tissue aging; cancer progression;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP; 7 A; 9 C; 6 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       method related oligonucleotide
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Pred.
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                                                                                                                                                                       for
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No. 9.
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                                                                                                                                                                    investigating
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         risk of developing (early the presence/absence of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length
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                                                                                                                                                                      aging
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                                                                                                                                                                         tissue
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monitoring

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RESULT 312
ADL07545/c
ID ADL075
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AC ADL075
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AC ADL075
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OT 06-MAY
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AAH45828
ID AAH458
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Best Local S
Matches 24
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Best Local
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                                                                                                                                                          telomeres, involving hybridising a chromosomal DNA extracted from a sample and a labeled DNA probe with a sequence complementary to a repetitive telomeric sequence, and measuring the labeled signal of the hybridised probe to give the size of telomere. This can be used to investigate tissue aging and the progression of cancer and in monitoring the progness of patients. The present sequence is an oligonucleotide described in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   the prognosis of patients. The present sequence is an oligonucleotide described in the exemplification of the invention
 06-MAY-2004
                                                                                                                                                                                                                                                     Determining telomere size progression of cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAH45828;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAH45828 standard; DNA;
                  ADL07545;
                                  ADL07545 standard; DNA; 24 BP.
                                                                                                                                            Sequence
                                                                                                                                                                                                                    The present invention describes a method for determining the
                                                                                                                                                                                                                                     Disclosure; Page 5; 8pp; Japanese.
                                                                                                                                                                                                                                                                               WPI; 2001-360495/38.
                                                                                                                                                                                                                                                                                                                30-SEP-1999;
                                                                                                                                                                                                                                                                                                                               30-SEP-1999;
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                                                                                                                                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                 repetitive
                                                                                                                                                                                                                                                                                                                                                                                                         Telomere size determination; chromosomal DNA; probe; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                          Telomere size
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                                                                                                      Local 5.
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                                                                           ш
                                                                                                                   Similarity
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                                                                                          GCCTCCCAAAGTGCTGGGATTACA 405
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GCCTCCCAAAGTGCTGGGATTACA 1
                                                                           GCCTCCCAAAGTGCTGGGATTACA 24
                                                                                                                                                                                                                                                                                                                                                                                                 sequence; tissue
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2.4%;
llarity 100.0%;
Conservative
                                                                                                           Conservative 0;
                                                                                                                                           BP; 6 A;
 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                         determination method related oligonucleotide #1.
                                                                                                                                                                                                                                                                                                                 99JP-00279948
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                                                                                                                                           7 C; 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    6 C; 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           24 BP
                                                                                                                                                                                                                                                            useful for investigating aging
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                                                                                                                                           G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                 aging;
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Pred. No.
                                                                                                                   Score 24;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    6 T; 0 U;
                                                                                                            Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                  cancer
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                                                                                                                   DB 1; Le 8.2e+02;
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hes 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                progression;
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                                                                                                                          Length 24;
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RESULT 313
AAH91303/c
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a polypeptide-Sec24 protein-31.35 and the polypeptide encoding it. Also included are the process for preparing the polypeptide by recombinant DNA technology, the application of the polypeptide in treating diseases such as cancer and HIV infection, the antagonist against the polypeptide (and its therapeutic action) and the application of the polypeptide encoding this polypeptide. The present sequence is an RT-(reverse transcriptase) PCR primer used to isolate cDNA
                                                                                                                                                                                                                                                  Human; inflammatory bowel disease; Crohn's disease; ulcerative colitis; single nucleotide polymorphism; SNP; chromosome 19p13; paternity test;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 24 BP; 4 A; 9 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               29-JAN-2003.
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                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                   chromosome
                                                                                                                                                                                                                                                                                                                    Human inflammatory bowel disease associated polymorphic site #378
                                                                                                                                                                                                                                                                                                                                                              09-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                        AAH91303;
                                                                                                                                                                                                                                                                                                                                                                                                                                                AAH91303 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 3; SEQ ID NO 3; 31pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Polypeptide-Sec24 protein-31.35.
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14-JUN-2001.
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                                                                                                                       misc_feature
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ilarity 100.0%;
Conservative (
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                                                                                                                           Location/Qualifiers
                                                                                                       *tag=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     protein-31.35.
                                                                                                                                                                                                                                 forensic test; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PCR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DEV INC
                                                                                 "SNP, optionally A or G at this position"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cancer; HIV infection; PCR; RT-PCR;
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11-DEC-2000; 2000WO-US033632.

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RESULT 314
AAA03956
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Best Loc
Matches
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 28 BP; 7 A; 5 C; 9 G; 6 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              polymorphisms associated with inflammatory bowel diseases such as ulcerative colitis and Crohn's disease. The methods can be used to dete the presence of genetic golymorphisms associated with inflammatory bowe disease and correlating their occurrence with disease states. They may used in this way for phenotypic correlations, forensics, paternity testing, medicine and genetic analysis. The present sequence is a polymorphic site described in the exemplification of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Testing for the presence of polymorphisms associated with inflammatory bowel disease, using a hybridization assay.
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10-APR-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention describes a method for detecting the presence of
                                                                                                                                                                                                                                                                                                                Polymorphism; hypertension; agammaglobulinemia; diabetes insipidus;
Lesch-Nyhan syndrome; muscular dystrophy; Wiskott-Aldrich syndrome;
Fabrys disease; familial hypercholesterolemia; hereditary spherocytosis;
                                                                                                                          07-MAY-1998;
03-MAY-1999;
                                                                                                                                                                                                                  EP955382-A2
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            Novel nucleic acids containing polymorphisms used
                                     WPI; 2000-107928/10
                                                                                                                                                               07-MAY-1999;
                                                                                                                                                                                        10-NOV-1999
                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                    Ehlers-Danlos
                                                                                                                                                                                                                                                                               familial colonic polyposis; osteogenesis
                                                                                                                                                                                                                                                                                           tuberous sclerosis; hereditary hemorrhagica telangiectasia;
                                                                                                                                                                                                                                                                                                        polycystic kidney disease; von Willebrands disease; forensic; human;
                                                                                                                                                                                                                                                                                                                                                                      Polymorphic
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                                                                                     AFFYMETRIX INC.
UNIV CASE WESTI
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                                                             Chakravarti
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                                                                                                                                                                                                                                                                                                                                                                      fragment of hypertension associated gene
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2000US-0196046P.
                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                       CASE WESTERN RESERVE.
                                                                                                                          98US-0084641P.
99US-00304232.
                                                                                                                                                               99EP-00250150
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                                                               Haluska MK;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  23.8;
No. 9
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                                                                                                                                                                                                                                                                                  imperfecta;
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             the
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              diagnosis
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Claim 1; Page
21; 53pp; English
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The invention provides polymorphic fragments of genes associated with CC hypertension. The nucleic acids including the polymorphic sites can be CC used as probes or primers for expressing variant proteins. Detection of CC the polymorphisms is useful in designing prophylactic and therapeutic CC regimes customized to underlying abnormalities. The polymorphisms can be CC used for association studies for hypertension, and in hypertension with CC diagnostic assays. Where the polymorphisms have strong correlation with CC hypertension. This information can be used to find the precise role in hypertension. This information can be used to find the precise role of a CC polymorphism in the disease, and this can be used to identify potential CC drugs which combat the disease e.g. agammaglobulinemia, diabetes consistion with other disease e.g. agammaglobulinemia, diabetes consistion with other diseases e.g. agammaglobulinemia, diabetes consistion, Lesch-Nyhan syndrome, muscular dystrophy, Wiskott-Aldrich CC syndrome, Fabrys disease, familial hypercholesterolemia, polycystic chidney disease, hereditary spherocytosis, von Willebrands disease, colonic polyposis, Ehlers-Danlos syndrome, osteogenesis imperfecta, and cute intermittent porphyria. The polymorphic forms can also be used in forensics to identify individuals

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Sequence 29
BP; 7 A; 4 C; 6 G; 11 T; 0 U; 1 Other;
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Query Match Best Local (Matches Local Similarity 1073 TIGTATTTCATTAGAGGCGGGGTTTCAC 1101 \vdash TTGTATTTTCAGTAKAGACAGGGTTTCAC Conservative 2.4%; Score 23.8; µb .. Mismatches 29 DB 1; Length Indels 29; , , Gaps 0

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AAA04662;
                  AAA04662 standard;
                   DNA;
                    29
                   BP.
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22-MAY-2000

(first entry)

밁 S

RESULT 315
AAA04662
XX AAA046
XX AAA046
XX Polymo
XX Famili
XX EP9553
XX EP9553
XX A CUYCAXX O7-MAY
XX O7-MAY
XX O7-MAY
XX O7-MAY
XX (AFFYPA (UYCAXX (Polymorphism; hypertension; agammaglobulinemia; diabetes insipidus; Lesch-Nyhan syndrome; muscular dystrophy; Wiskott-Aldrich syndrome; Fabrys disease; familial hypercholesterolemia; hereditary spherocytosis; polycystic kidney disease; von Willebrands disease; forensic; human; tuberous sclerosis; hereditary hemorrhagica telangiectasia; familial colonic polyposis; osteogenesis imperfecta; porphyria; Polymorphic Ehlers-Danlos fragment of hypertension associated gene TBXA2R. syndrome; 88.

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Homo sapiens
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EP955382-A2

10-NOV-1999.

07-MAY-1999; 99EP-00250150

03-MAY-1999; 07-MAY-1998; 98US-0084641P. 99US-00304232.

(AFFY-) (UYCA-) AFFYMETRIX INC. CASE WESTERN RESERVE.

'n, Chakravarti Þ Haluska MK

WPI; 2000-107928/10

hypertension.

Novel nucleic acids containing polymorphisms used in the diagnosis hypertension of.

Claim 1;

43; 53pp; English.

invention provides polymorphic fragments of genes associated with ertension. The nucleic acids including the polymorphic sites can be

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RESULT 316
AAA04486/c
       PRINTER PRINTE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity
Matches 25; Conservat
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           polycystic kidney disease, tuberous sclerosis, heredit familial colonic polyposis.
Claim 1; Page 38; 53pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                 07-MAY-1998;
03-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Polymorphic fragment of hypertension associated gene PGIS
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                                                                                                      Novel nucleic acids containing polymorphisms used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   07-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10-NOV-1999
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                                                                                                                                                                                                                                                                                                              AFFYMETRIX INC.
UNIV CASE WESTERN RESERVE.
                                                                                                                                                                                                                                           Chakravarti A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           colonic polyposis; osteogenesis imperfecta; porphyria;
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ilarity 86.2%;
Conservative
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99US-00304232.
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Pred. No. 9
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07-MAY-1998; 03-MAY-1999;

98US-0084641P. 99US-00304232.

Claim 1; Page 18; 53pp;

hypertension.

Fan JB,

Chakravarti A,

Haluska MK

(AFFY-) AFFYMETRIX INC. (UYCA-) UNIV CASE WESTERN RESERVE.

Novel nucleic acids containing polymorphisms used in the diagnosis

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RESULT 317
AAA03878/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention provides polymorphic fragments of genes associated with CC hypertension. The nucleic acids including the polymorphic sites can be CC used as probes or primers for expressing variant proteins. Detection of CC the polymorphisms is useful in designing prophylactic and therapeutic regimes customized to underlying abnormalities. The polymorphisms can be CC used for association studies for hypertension, and in hypertension CC diagnostic assays. Where the polymorphisms have strong correlation with CC hypertension, within a gene, they are likely to have a causative role in CC hypertension. This information can be used to find the precise role in CC polymorphism in the disease, and this can be used to identify potential CC drugs which combat the disease. The polymorphisms can be tested for CC association with other diseases e.g. agammaglobulinemia, diabettes CC insipidus, Lesch-Nyhan syndrome, muscular dystrophy, Wiskott-Aldrich CC syndrome, Fabrys disease, familial hypercholesterolemia, polycystic kidney disease, hereditary spherocytosis, von Willebrands disease, familial CC colonic polyposis, Ehlers-Danlos syndrome, osteogenessis imperfecta, and CC acute intermittent porphyria. The polymorphic forms can also be used in CC forensics to identify individuals
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Polymorphism; hypertension; agammaglobulinemia; diabetes insipidus; Lesch-Nyhan syndrome; muscular dystrophy; Wiskott-Aldrich syndrome; Fabrys disease; familial hypercholesterrolemia; hereditary spherocytosis; polycystic kidney disease; von Willebrands disease; forensic; human; tuberous sclerosis; hereditary hemorrhagica telangiectasia; familial colonic polyposis; osteogenesis imperfecta; porphyria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 29 BP; 9 A; 10 C; 4 G; 5 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAA03878 standard;
                                                                                                                                                                                                                                                                                                07-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                               Ehlers-Danlos
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Polymorphic fragment of hypertension associated gene AE1.
                                                                                                                                                                                                                                                                                                                                     10-NOV-1999
                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          29
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TTAGTAGAGATGGAGTTTCTCCATGTTGG 205
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TTAGTAGAGACGGGRTTTCGCCATGTTGG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                 syndrome;
                                                                                                                                                                                                                                                                                                99EP-00250150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 23.8; DB 1;
Pred. No. 9.5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 29;
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CC The invention provides polymorphic fragments of genes associated with CC hypertension. The nucleic acids including the polymorphic sites can be CC used as probes or primers for expressing variant proteins. Detection of CC the polymorphisms is useful in designing prophylactic and therapeutic CC regimes customized to underlying abnormalities. The polymorphisms can be CC used for association studies for hypertension, and in hypertension of CC diagnostic assays. Where the polymorphisms have strong correlation with hypertension. This information can be used to find the precise role in CC polymorphism in the disease, and this can be used to identify potential CC drugs which combat the disease. The polymorphisms can be tested for CC insipidus, Lesch-Nyhan syndrome, muscular dystrophy, Wiskott-Aldrich CC syndrome, Fabrys disease, familial hypercholesterolenia, polycystic cc kidney disease, hereditary spherocytosis, von Willebrands disease, familial colonic polyposis, Ehlers-Danlos syndrome, softeogenesis imperfecta, and acute intermittent porphyria. The polymorphic forms can also be used in forensics to identify individuals
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 318
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 29 BP; 7 A; 6 C; 11 G; 4 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                       07-MAY-1998;
03-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                         07-MAY-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EP955382-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        familial colonic polyposis; osteogenesis imperfecta; porphyria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   polycystic kidney disease; von Willebrands disease; forensic; human; tuberous sclerosis; hereditary hemorrhagica telangiectasia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Lesch-Nyhan syndrome; muscular dystrophy; Wiskott-Aldrich syndrome;
Fabrys disease; familial hypercholesterolemia; hereditary spherocytosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Polymorphism; hypertension; agammaglobulinemia; diabetes insipidus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             22-MAY-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAA04660 standard;
Claim 1; Page 43; 53pp;
                                                                                                                                              WPI; 2000-107928/10.
                                                                                     Novel nucleic acids containing polymorphisms used in the
                                                                                                                                                                                                                                                                                          (AFFY-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    670 TIGGCICACTGCAACCICTGCCTCCCGGG 698
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                                                                                                                                                                                                                                                                                          AFFYMETRIX INC
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                                                                                                                                                                                                        Chakravarti A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   fragment of hypertension associated gene TBXA2R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                  CASE WESTERN RESERVE
                                                                                                                                                                                                                                                                                                                                                 98US-0084641P.
99US-00304232.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29
      English
                                                                                                                                                                                                           Haluska MK;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; Score 23.8; DI
; Pred. No. 9.5e.
1; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                           diagnosis
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The invention provides polymorphic fragments of genes associated with

The invention provides polymorphic fragments of genes associated with hypertension. The nucleic acids including the polymorphic sites can be

Claim 1; Page

38; 53pp; English.

Novel nucleic acids containing polymorphisms used

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diagnosis of

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RESULT 319
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Lesch-Nyhan syndrome; muscular dystrophy; Wiskott-Aldrich syndrome; Fabrys disease; familial hypercholesterolemia; hereditary spherocytosis; polycystic kidney disease; von Willebrands disease; forensic; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                         polycystic kidney disease; von Willebrands disease; forensic; human; tuberous sclerosis; hereditary hemorrhagica telangiectasia; familial colonic polyposis; osteogenesis imperfecta; porphyria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 29 BP; 4 A; 11 C; 8
                                                                                                                                                                                                                                             07-MAY-1998;
03-MAY-1999;
                                                                                                                                                                                                                                                                                             07-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Polymorphic fragment of hypertension associated gene PGIS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAA04502 standard; DNA; 29
                                                                                                                            WPI; 2000-107928/10
                                                                                                                                                                                                                                                                                                                                                                                                                               Ehlers-Danlos
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Polymorphism; hypertension; agammaglobulinemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             22-MAY-2000
                                                                                                                                                                                                                                                                                                                            10-NOV-1999
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                                                                                                                                                             JB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          876 GGCGTGAGCCACCACGCCCGGCTTATTTT 904
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                                                                                                                                                                                           AFFYMETRIX INC.
UNIV CASE WESTERN RESERVE.
                                                                                                                                                             Chakravarti
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
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99US-00304232.
                                                                                                                                                                                                                                                                                             99EP-00250150
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                                                                                                                                                             Haluska MK;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 23.8;
Pred. No. 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               29
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1; Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  diabetes insipidus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
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RESULT 320
AAA04661
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Best Local Similarity
Matches 25; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Polymorphism; hypertension; agammaglobulinemia; diabetes insipidus;
Lesch-Nyhan syndrome; muscular dystrophy; Wiskott-Aldrich syndrome;
Fabrys disease; familial hypercholesterolemia; hereditary spherocytosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
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Claim 1; Page 43;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      07-MAY-1998;
03-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            07-MAY-1999;
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tuberous sclerosis; hereditary hemorrhagica telangiectasia;
familial colonic polyposis; osteogenesis imperfecta; porphyri;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22-MAY-2000
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                                                                                                                                                                                                                                                                  Fan JB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1034
                                                                                                                                                                                        2000-107928/10
                                                                                                         nucleic acids
                                                                                                                                                                                                                                                                                                                                             AFFYMETRIX INC.
UNIV CASE WESTERN RESERVE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CTGGGATTACGGGCACCTGCCACACC 1062
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                                                                                                                                                                                                                                                              Chakravarti A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                fragment
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      98US-0084641P
99US-00304232
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             polyposis; osteogenesis
53pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA;
                                                                                                         containing polymorphisms used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    of hypertension associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   29
                                                                                                                                                                                                                                                                  Haluska MK;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ВP
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Pred. No. 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             imperfecta; porphyria;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TBXA2R.
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The invention provides polymorphic fragments of genes associated with hypertension. The nucleic acids including the polymorphic sites can be used as probes or primers for expressing variant proteins. Detection o

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The invention provides polymorphic fragments of genes associated hypertension. The nucleic acids including the polymorphic sites cused as probes or primers for expressing variant proteins. Detect the polymorphisms is useful in designing prophylactic and therape

Detection of

can be

1; Page 32;

53pp; English

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RESULT 321
AAA04307
ID AAA043
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                the polymorphisms is useful in designing prophylactic and therapeutic regimes customized to underlying abnormalities. The polymorphisms can be used for association studies for hypertension, and in hypertension diagnostic assays. Where the polymorphisms have strong correlation with hypertension. This information can be used to find the precise role in hypertension in the disease, and this can be used to identify potential drugs which combat the disease. The polymorphisms can be tested for association with other disease e.g. agamaglobulinemia, diabetes insipidus, Lesch-Nyhan syndrome, muscular dystrophy, Wiskott-Aldrich syndrome, Fabrys disease, familial hypercholesterolemia, polycystic kidney disease, hereditary spherocytosis, von Willebrands disease, tuberchytosis, ron Willebrands disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                            Polymorphism; hypertension; agammaglobulinemia; diabetes insipidus; Lesch Nyhan syndrome; muscular dystrophy; Wiskott-Aldrich syndrome; Fabrys disease; familial hypercholesterolemia; hereditary spherocyt polycystic kidney disease; von Willebrands disease; forensic; human tuberous sclerosis; hereditary hemorrhagica telangiectasia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            tuberous sclerosis, hereditary hemorrhagica telangiectasia, familial colonic polyposis, Ehlers-Danlos syndrome, osteogenesis imperfecta, acute intermittent porphyria. The polymorphic forms can also be used forensics to identify individuals
                                                                                                                                                                                                                                                  07-MAY-1998;
03-MAY-1999;
                                                                                                                                                                                                                                                                                             07-MAY-1999;
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                                                                                                                                                                                                                                                                                                                                                        EP955382-A2
                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
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                                                                                                  hypertension.
                                                                                                                                             WPI; 2000-107928/10.
                                                                                                                                                                          Fan JB,
                                                                                                                                                                                                                                                                                                                                                                                                                              familial colonic polyposis; osteogenesis imperfecta; porphyria;
                                                                                                                                                                                                       (UYCA-)
                                                                                                                                                                                                                       (AFFY-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   650 TGGAGTGCAGTGGCGCAATCTTGGCTCAC
                                                                                                                nucleic acids containing polymorphisms used in the diagnosis
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UNIV CASE WEST
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                                                                                                                                                                          Chakravarti A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     fragment of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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99US-00304232.
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                                                                                                                                                                                                         WESTERN RESERVE.
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Pred. No. 9.5e
1; Mismatches
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RESULT 322
AAZ09548/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CC regimes customized to underlying abnormalities. The polymorphisms can be considered to underlying abnormalities. The polymorphisms can be considered to assays. Where the polymorphisms have strong correlation with the hypertension, within a gene, they are likely to have a causative role in the polymorphisms can be used to find the precise role of a compartension. This information can be used to identify potential confunction with disease, and this can be used to identify potential consistency which combat the disease. The polymorphisms can be tested for consipidus, Lesch-Nyhan syndrome, muscular dystrophy, Wiskott-Aldrich consistency disease, hereditary spherocytosis, von Willebrands disease, tuberous sclerosis, hereditary spherocytosis, von Willebrands disease, colonic polyposis, Ehlers-Danlos syndrome, coteogenesis imperfecta, and coute intermittent porphyria. The polymorphic forms can also be used in corresponds to identify individuals
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Best Local S
Matches 25
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                                                                                                                                                                                                                                                                 12-FEB-1998;
30-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Apo E; Apo B; hyperlipidemia; human; treatment; hepatocyte; apoprotein; Apo A1; low density lipoprotein; LDL; blood; therapy; atherosclerosis; high density lipoprotein; HDL; cholesterol; coronary heart disease; high density lipoprotein; HDL; cholesterol; coronary heart disease; Alzheimer's disease; hypobetalipoproteinemia; dysbetalipoproteinemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      08-NOV-1999
                                                                                                                                    Mutating apolipoprotein
                                                                                                                                                                 WPI; 1999-527333/44
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98US-00108006.
                                                                                                                                                                                                                                                                                                            98WO-US017908
                                                                                                                                                                                             Ħ,
                                                                                                                       protein genes in hepatocytes to control cholesterol treating or preventing hyperlipidemia, particularl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     entry)
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                                                                                                                                                                                             Bandyopadhyay PT,
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                                                                                                                                                                                             Roy-Chowdhury J;
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This invention describes a novel method for the genetic treatment hyperlipidemia by altering genes, in hepatocytes, for apoprotein E or Al. Low density lipoprotein (LDL) levels in the blood are recaltering an apo B gene (I) in a hepatocyte. The invention describe method for the therapeutic and/or prophylactic method involving al

describes are reduced

(apo) o H

A method has been developed for determining a patient's susceptibility t an inflammatory disorder. The method comprises the detection of an interleukin 1 (II-1) (44112332) haplotype in a sample obtained from the patient, where its presence indicates susceptibility to an inflammatory disorder. AAX16607 to AAX16631 represent PCR primer used in the method for detecting the II-1 (44112332) haplotype. The method provides kits for the early prediction of a patient's susceptibility to inflammatory artery disease, osteoporosis, nephropathy disorders, including coronary artery disease, osteoporosis, nephropathy

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49pp; English

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ribes a galtering

diabetes mellitus,

Claim 31; Page 51;

106pp; English

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RESULT 323
AAX16609
ID AAX166
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밁
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Graves disease; subjects to the control of the c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  an apo E gene in hepatocytes by introducing the mutations Arg112Cys, Arg158Cys or Cys158Arg and a method for ameliorating atherosclerosis by Algering the apo Al gene in a hepatocyte so that the altered protein can dimerize. Altering expression of apo genes regulates levels of high and low density lipoprotein cholesterol. Altering expression of apo B. E and Al genes is used to treat or prevent atherosclerosis, coronary heart disease, Alzheimer's disease, hypobetalipoproteinemia, and dysbetalipoproteinemia. AAZ09545-Z09548 represent primers used in the manipulation of the human Apo E protein described in the method of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 25
                                                                                                                                                                                                                                                                                                   New method of determining a patient's susceptibility to inflammatory disorders - by detecting the presence of an IL-1 (4411232) haplotype, useful in designing treatment strategies that modulate the activity of proteins produced by the IL-1 gene cluster.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (DUFF/)
(COXA/)
(CAMP/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Duff G,
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COX A.
CAMP N J.
DE GIOVINE 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cox A,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      IL-1; haplotype; inflammatory disorder; alopecia
y disease; osteoporosis; nephropathy; diabetes mel
; systemic lupus erythamatosus; lichen sclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   97GB-00011040
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9e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human, single-nucleotide polymorphism; SNP; SNAP23; ss; PCR primer; synaptosome associated protein of 23 kilodaltons; diabetes; obesity; cardiovascular disorder; hypertension; cancer; drug screening; forensic medicine; paternity testing.
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                                                                                                                                                                                                                                                                                                                                                                                                                       Polymorphic sites useful for the diagnosis of metabolic diseases involving glucose homeostasis e.g. diabetes, obesity and cardiovascular
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The invention relates to a nucleic acid between 10 and 100 bases comprising at least 10 contiguous nucleotides including a polymorphic site or an adjacent base, from 6 polymorphisms from 5 human genes, RGS5 (regulator of G-protein signalling 5), SNAP23 (synaptosome associated protein of 23 kilodaltons), ALDOB (aldolase B), GOS8/RGS2 (helix-loop-helix basic phosphoprotein) and PPPICB (not defined). The invention also nucleic acid comprising but polymorphisms in a method comprising analysing a nucleic acid comprising by obtaining nucleic acid samples from individuals, determining a base occupying any one of the polymorphic or other sites in linkage disequilibrium with them and testing each

tested for

Disclosure; Page 8; 41pp; English.

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AAS14581
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          The invention relates to a nucleic acid between 10 and 100 bases CC comprising at least 10 contiguous nucleotides including a polymorphic comprising at least 10 contiguous nucleotides including a polymorphic cc site or an adjacent base, from 6 polymorphisms from 5 human genes, RGS5 CC (regulator of G-protein signalling 5), SNAP23 (synaptosome associated protein of 23 kilodaltons), ALDOB (aldolase B), COSB/RGS2 (helix-loop-cc helix basic phosphoprotein) and PPPICB (not defined). The invention also CC relates to using such polymorphisms in a method comprising analysing a nucleic acid comprising by obtaining nucleic acid samples from CC individuals, determining a base occupying any one of the polymorphic or CC individuals, determining a base occupying any one of the polymorphic or CC individual for the presence of a phenotype and correlating the presence CC individual for the presence of a phenotype and correlating the presence CC individual for the presence of a phenotype and correlating the presence CC determined is used as a diagnostic tool for diseases such as diabetes, CC cancers. The polymorphisms may also be used for drug screening, with other contractive traces and could be resented for association with other contractive traces and could be resented for association with other contractive traces and could be resented for association with other contractive traces and could be resented.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ma Y,
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                                                                                                                                                                                                                                                                                                                                                                                           disorders
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                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 8; 41pp; English.
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polymorphisms may also be u
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RESULT 326
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SNAP23
                                                                                         The present invention relates to a new oligonucleotide label-domain comprising the sequence (CTATTTT) n and its complement (AAAATAG) n, where n is 1. The probe sets of the invention are useful for detecting kappa or lambda-immunoglobulin light chain mRNA or corresponding heteronuclear RNA, CMV (cytomegalovirus) immediate early RNA, EBV (Epstein-Barr virus) early RNA 1 and RNA 2, and human Alu repetitive satellite genomic sequences. The invention is a useful generic sequence for incorporation into oligonucleotide probes for detecting gene-specific sequences within cells or tissue samples in in situ hybridisation analysis and for attaching a label to immunoglobulins or other proteins for detecting haptens and antigens in immunohistochemical analyses. The present nucleic acid sequence represents one of a collection (ABK70376-ABK70501) of oligonucleotide probes that were used in the invention for detecting or localising a plurality nucleic acid target gene or antigen within a cell
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within
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                                                                                                                                                                                                                                                                                                                                                     Novel oligonucleotide label-domain for incorporation into oligonucleotide probes useful for detecting or localizing nucleic acid target genes
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ı a cell
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24; Conserv
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                                                       BP; 4
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                                                                                                                                                                                                                                                                                                                                                                                                                          Connaughton J;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA;
           2.4%;
96.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       7 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               t sequence represents a PCR primer used to amplify a SNP (single-nucleotide polymorphism) from human
                                                                                                                                                                                                                                                                                                         71pp;
                                                       5 C; 13
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Pred. No. 9
Score 23.4; DB 1;
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                      The present invention relates to methods for identifying a test substance CC that modulate the immune response in a genotype specific manner. Methods CC of the invention involve genotyping subjects to identify those having a CC genotype (e.g. interleukin-1; IL-1) associated with one or more CC inflammatory disorder. The method comprises genotyping a subject having a CC an inflammatory disease-associated genotype and observing a biomarker in CC the subject before and after the subject is contacted with the test CC substance. The methods or cells associated with inflammatory diseases are CC aspecific biological response in subjects having inflammatory diseases are CC associated genotype, where the genotype is associated a pre-disposition to one or more of periodontal disease, coronary artery disease, CC Alzheimer's disease, atherosclerosis, osteoporosis, insulin-dependent CC diabetes, diabetic retinopathy, end-stage renal disease, diabetic retinopathy, end-stage renal disease, diabetic CC nephropathy, hepatic fibrosis, alopecia areata disease, diabetic contacted arthritis, juvenile chronic arthritis, complete sclerosis, idiopathic pulmonary fibrosis, sepsis and acne. The CC multiple sclerosis, idiopathic pulmonary fibrosis, sepsis and acne. The CC multiple sclerosis, idiopathic pulmonary fibrosis, sepsis and acne. The complication of the invention for cenotyping IL-1A (925/926) gene. This primer is used in the second of the invention for cenotyping IL-1A (925/926) gene. This primer is used in the second of the invention for cenotyping IL-1A (925/926) gene.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Screening a substance in a subject for modulating an immune response, comprises genotyping to identify the test subject, and observing a biomarker before and after contacting the subject with the test
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PCR primer #1, used for genotyping human IL-1A (gz5/gz6) marker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAD27391 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example; Page 42;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               nootropic;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (INTE-) INTERLEUKIN GENETICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          periodontal disease; Alzheimer's disease; atherosclerosis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     _
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RESULT 328
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ7; MDZ12; chromosome 7g22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                   proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is encoded at chromosome 7g22.1, MDZ4 is encoded at chromosome 6p21.3-22.2, MDZ7 is encoded at chromosome 1fp11.2 and MDZ12 is encoded at chromosome 1fp11.2 and MDZ12 is encoded at chromosome 15g26.1. The MDZ3, MDZ4, MDZ7, and MDZ12 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12 genetic locus. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADB04743 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Shannon M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                05-FEB-2003
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                                                                                                                                                                                                                                                                                                       The present invention relates to novel human zinc finger-containing
                                                                                                                                                                                                                                                                                                                                     Example 8;
                                                                                                                                                                                                                                                                                                                                                               New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ4, MDZ7 or MDZ12, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  02-AUG-2001; 2001US-00922181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                30-JUL-2002; 2002EP-00016874
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        developmental
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              Local Similarity
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                                                                                        The present sequence was used to illustrate
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              2.4%;
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              Score 23.4;
Pred. No. 96
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3 T; 0 U;
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   Mismatches
                                                                                                                                                                                                                                                                                                                                   English.
                                                           U; 0 Other;
               9e+02;
                            DB 1; Length
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RESULT 329
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RESULT 330
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                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                          The present invention provides a method for determining the predisposition of a subject to early-onset menopause (EOM). The invention is useful in diagnosing, treating and preventing early-onset menopause. The present sequence is human secreted interleukin-1A (IL-1A) amplifying PCR primer. The sequence is used in exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Early-onset menopause; EOM; diagnosis; therapy; human; interleukin-1A; II-1A; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human interleukin-1A (gz5/gz6)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADN48862 standard;
                                                                                                                                                                                                                                                                                       Sequence 25 BP; 5 A; 7 C; 10 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                              Determining the predisposition to early-onset detecting in the subject interleukin (IL)-1RN
                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2004-354679/33.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  04-AUG-2000; 2000US-00632657.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       04-MAY-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US6730476-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15-JUL-2004
                                                                                                       01-FEB-2000
                                                                                                                           AAZ37279;
                                                                                                                                               AAZ37279 standard;
                                                                                                                                                                                                                                                                                                                                                                          Example;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
                  Homo sapiens
                              Synthetic
                                                                                  PCR primer for SGRF coding sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (INTE-) INTERLEUKIN GENETICS INC.
                                                   mmune
                                                                                                                                                                                                                                                        Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                    ຸດ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          770
                                                                                                                                                                                                                        867 GGGATTACAGGCGTGAGCCACCACG
                                                              human;
                                                                                                                                                                                                   ب
                                                                                                                                                                                                                                              24;
                                                  system;
                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TTTTGTATTTTTAGTAGAGATGGGG
                                                                                                                                                                                                                                                                                                                                                                           SEQ ID NO 25;
                                                                                                                                                                                                   GGGATTACAGGCGTGAGCCACCGCG
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Kornman K,
                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry
                                                             Interleukin-6 G-CSF related factor; cell proliferation;
                                                                                                       (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             99US-00345217
                                                   haematopoietic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA;
                                                                                                                                                DNA;
                                                                                                      entry)
                                                                                                                                                                                                                                                        2.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Van Dijk S;
                                                                                                                                                                                                                                                                                                                                                                         57pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25
                                                                                                                                                 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP
                                                                                                                                                ВP
                                                                                                                                                                                                                                             <u>,</u>
                                                                                                                                                                                                                                                        Score 23.4;
Pred. No. 9e
                                                   system; therapy; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  amplifying PCR primer
                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            794
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25
                                                                                                                                                                                                                          891
                                                                                                                                                                                                    25
                                                                                                                                                                                                                                                                  DB 1;
                                                                                                                                                                                                                                                                                                                                                                                               menopause comprises (+2018) allele 2.
                                                                                                                                                                                                                                                                   Length
                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      #
                                                                                                                                                                                                                                              0,
                                                                                                                                                                                                                                              Gaps
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WO9954357-A1

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RESULT 331
AAH38611/c
ID AAH386
XX AAH386
AC AAH386
AC SNP sp
XX Single
CW Single
KW Single
KW Single
KW Solycy
KW acute
KW acute
KW inflam
XX Homo s
XX WO2001
XX GE-APR
XX GORCH-
PH 15-OCT
XX ORCH-
PH NEW 96
PT Acid s
XX New 96
PT Acid s
XX AAH386
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Best Local S
Matches 24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28-OCT-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This sequence represents a PCR primer used to isolate DNA encoding the Interleukin-6 G-CSF related factor (SGRF) protein of the invention. The protein as member of the II-6/G-CSF/MSF family. The protein can be used in drugs for treating diseases due to abnormality of cell proliferation in the immune system and haematopoietic system
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2000-013230/01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14-APR-1998;
                                                                                                                                                                                                                                                                                                                                                       Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel cytokine-like protein, with activity of supporting proliferation myeloid cells, useful in treating abnormality of cell proliferation in
                                                                                                                                                                                                                                                                                                        polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis; inflammation; forensic investigation; paternity analysis; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 5; Page 23; 69pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (CHUS ) CHUGAI RES INST MOLECULAR MEDICINE INC
               New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nucacid sample.
                                                                                                                                                                                           13-OCT-2000; 2000WO-US028436
                                                                                                                                                                                                                        26-APR-2001
                                                                                                                                                                                                                                                     WO200129262-A2
                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                   SNP specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                 14-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAH38611
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAH38611 standard; DNA; 27
                                                                                                                                                             15-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        immune and hematopoiesis
                                                                           WPI; 2001-290930/30
                                                                                                                                   (ORCH-) ORCHID BIOSCIENCES INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                537
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CCTGCCTCAGCCTCCCAAGCAGCTG 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CCTGCCTCAGCCTCCCAAGTAGCTG 561
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP; 4 A; 5 C; 12 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                     SNPE primer SEQ ID 1407.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        98JP-00121805
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       99WO-JP001997
                                                                                                                                                               99US-0160096P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           96.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         systems.
                                                                                                       Z
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 23.4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             .4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 27;
                              in a nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0,
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RESULT 332
AAA27185
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cc primer extension (SNPE) primers, and the sequences of regions flanking concludes kits for determining the presence or absence of a SNP, using the coligonucleotides of the invention. The PCR primers are used to amplify a coligonucleotides of the invention. The pCR primers are used to amplify a coligonucleotides are useful for genotyping a nucleic acid sample by performing a single-nucleotide primer extension reaction. The CC assess by association analysis the genotype of an individual or group of cindividuals, having a pathological phenotypic trait suspected of being caused by one or more SNPs. Phenotypic traits include diseases e.g. cagammaglobulinaemia, diabetes insiplous, Lesch-Nyhan syndrome, muscular cystrophy, familial hypercholesterolaemia, polycystic kidney disease, costeogenesis imperfecta and acute intermittent porphyria. Phenotypic traits also include symptoms of or susceptibility to multifactorial consists and include symptoms of or susceptibility to multifactorial confidence, including, rheumatoid arthritis, multiple sclerosis, microgranism. The method is also useful in forensic investigations and primer extension (SNPE) primer specific for a human SNP containing DNA comparer.
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                 P2; CX5C
allegic r
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1; Page 57; 83pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                 Reverse primer IL10 for target sequence human interleukin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 27 BP; 6
                          Measuring target polymucleotide sequences in biological samples by contacting sequence-selective primer pairs, forming conjugates with adaptor molecules, polymerizing target-identifier dimers and quantifying
                                                                                                                                                                                                                                                                                                                                                                                                                                                 11-SEP-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAA27185 standard; DNA; 28
                                                                                                                         Dolganov
                                                                                                                                                                                         16-NOV-1998;
                                                                                                                                                                                                                       12-NOV-1999;
                                                                                                                                                                                                                                                      25-MAY-2000.
                                                                                                                                                                                                                                                                                      WO200029621-A2
                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                         (GENE-)
                                                                                             2000-387825/33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           673 GCTCACTGCAACCTCTGCCTCCCGGGT 699
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               27
                                                                                                                                                                                                                                                                                                                                                                 Chemokine; Chromosome 5q31; gene therapy; asthma; PCR primer; rhinitis; urticaria; anaphylactic shock; hives; hay fever; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2.4%;
Similarity 88.9%;
                                                                                                                                                         GENELABS TECHNOLOGIES
                                                                                                                         ຸດ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GCTCACTGNAACCTCTGCCTCCGGGNT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                         Novikov
                                                                                                                                                                                         98US-00193320
                                                                                                                                                                                                                        99WO-US026931
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; 6 C; 10 G; 3 T; 0 U; 2 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 23.4;
Pred. No. 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               .4e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               27;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
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Disclosure; Page 100; 103pp; English

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RESULT 333
ACC84463
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CC A novel method for simultaneously determining the level of a number of control target polynucleotides in a sample has been disclosed. The method converted to the target levels in the varget sequence in direct correction to the target levels in the original sample. The target correction to the target levels in the original sample. The target corrected with either first or second adaptor sequences. The first and corrected with either first or second adaptor sequences. The first and corrected with either first or second adaptor sequences. The first and corrected with either first or second adaptor sequences. The first and correct through the target sequences. The adaptor sequences are then removed to leave target sequences in the adaptor sequences are then polymerised to form dimer multimers. The relative abundances of target identifiers in the multimer allow expression levels to be determined. This method is useful corrected under various conditions, stages of development and and disease tates, particularly where the target polymucleotide is present at low correctly. The method may also be used in the discovery and evaluation of addition to the method described above, the invention also includes the polymucleotide and polypeptide of P2. P2 is thought to be a member of a novel chemokine family, denoted CX5C and may be associated with immune function. Compositions of the P2 polymucleotide may be associated with immune correctly shock and conditions involving immune system hypersensitivity. The P2 polymucleotide to treat conditions using gene cutoff the passed to chromosome 5, within the cytokine gene cluster at 5q3l. The present sequence is the reverse primer conditions under the reverse primer and interleukin 10
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Best Local S
Matches 24
                                                                                                                                                                                                                                                                                                                      Unidentified
                                                                                                                                                                                                                                                                                                                                                       Cytostatic; Antibacterial; Immunosuppressive; Antiinflammatory; neural thread protein; NTP; tumour; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ACC84463 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                P-PSDB; ABR63258
                                 WPI; 2003-247999/24
                                                                      Averback
                                                                                                                                        19-JUL-2001; 2001US-0306150P.
19-JUL-2001; 2001US-0306161P.
16-NOV-2001; 2001US-0331477P.
                                                                                                                                                                                                             19-JUL-2002; 2002WO-CA001105
                                                                                                                                                                                                                                                  30-JAN-2003
                                                                                                                                                                                                                                                                                   WO2003008443-A2
                                                                                                                                                                                                                                                                                                                                                                                                             NTP peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                              28-AUG-2003
                                                                                                       (-OMYN)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           635 CTCTGTCACCCAGGCTGGAGTGCAG 659
                                                                                                       XOMYN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                    PA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CTCTGTCACCCAGGCTGGAGTACAG 28
                                                                                                                                                                                                                                                                                                                                                                                                           encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP; 5 A; 8 C; 8 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                         CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA;
                                                                                                                                                                                                                                                                                                                                                                                                         sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 23.4;
Pred. No. 9.
                                                                                                                                                                                                                                                                                                                                                                                                             #10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             9.7e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
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Gaps

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The present invention relates to a neural thread protein (NTP) peptide referred to as cell death peptide. Thought to be cytostatic, antibacterial, immunosuppressive and antilinflammatory. It is useful for treating a condition in a patient requiring removal or destruction of cells, for treating a condition such as benign or malignant tumor, inflammatory disease, autoimmune disease and infectious disease. The peptide useful for treatment is derived from the amino acid sequence for a pancreatic thread protein. The peptide is conjugated, linked or bound to a molecule chosen from antibody or its fragment, antibody-like binding molecule, where the molecule has a higher affinity for binding to a tumor or other target than binding to other cells. Treatment using NTP peptides can remove benign tumors with less risk and fewer of the undesirable side effects of surgery. The present sequence is an NTP encoding sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel neural thread protein peptide, referred as cell death peptide, useful for treating prostatic hyperplasia, psoriasis, eczema, dermatos atherosclerosis, cosmetic modification to skin, throat, mouth, muscle.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 17; 77pp; English.
Sequence 28
   BP; 6 A; 9 C; 9
   G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         dermatosis,
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RESULT 334
                                                             Query Match
Best Local S
Matches 24
                                      725 CCTGAGTAGCTGGGACTACAGGCGC 749
                                                               24;
                            4 CCAGAGTAGCTGGGACTACAGGCGC
                                                                       Similarity
                                                               Conservative
                                                                     2.4%;
                                                              0,
                                                                      Score 23.4; DB 1, Pred. No. 9.7e+02
                                                               Mismatches
                             28
                                                                                 ۲.
                                                               ۲,
                                                                                Length
                                                               Indels
                                                               0,
                                                               Gaps
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LRRC8; B-cell surface membrane; immunosuppressive; non-gamma globulin blood disease; maturation; human; ss; chromosome 9; chromosome 20; translocation.
                                                                                                                                                                                                                                RT-PCR primer 2 related to human blood disease-related LRRC8 mutant.
                                                                                                                                                                                                                                                             ADP70455;
                                                                                                                                                             20-MAY-2004
                                                                                                                                                                           JP2004141048-A.
                                                                                                                                                                                        Homo sapiens.
                                                                                                                                                                                                                                               12-AUG-2004
                                                                                                                                                                                                                                                                          ADP70455 standard; DNA; 28
                                                                                                                                                                                                                                               (first entry)
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sequence as given in the specification, or a receptor protein in which one or more amino acid residues are substituted, added or deleted, that expresses on the B-cell surface membrane of a non-gamma globulin diseased patient. The polypeptide of the invention demonstrates immunosuppressive activity and may be useful for screening a B-cell-associated disease

s to a novel LRRC8 protein comprising the specification, or a receptor prot

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fully defined in in which

therapeutic agent

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immunosuppressive

agent,

tor

congenital

Novel blood

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immunosuppressive agent.

LRRC8 protein useful as marker for congenital non-gamma globulin disease, or for screening B-cell associated disease therapeutic

WPI; 2004-382668/36

23-OCT-2002; 2002JP-00308855. 23-OCT-2002; 2002JP-00308855

(00SA-)

ZH OOSAKA SANGYO SHINKO OSAKA PREFECTURE.

Disclosure; SEQ ID NO 12; 46pp; Japanese

The invention relates

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ARAGULT 335
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                                                                                             The invention provides polymorphic fragments of genes associated with hypertension. The nucleic acids including the polymorphic sites can be used as probes or primers for expressing variant proteins. Detection of the polymorphisms is useful in designing prophylactic and therapeutic regimes customized to underlying abnormalities. The polymorphisms can be used for association studies for hypertension, and in hypertension diagnostic assays. Where the polymorphisms have strong correlation with hypertension, within a gene, they are likely to have a causative role in hypertension. This information can be used to find the precise role of a polymorphism in the disease, and this can be used to identify potential drugs which combat the disease. The polymorphisms can be tested for association with other disease e.g. agammaglobulinemia, diabetes insipidus, Lesch-Nyhan syndrome, muscular dystrophy, Wiskott-Aldrich annotrome. Fahrya disease familial hymeryheiderterolamia columniation of the proventical and the precise familial hymeryheiderterolamia.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         polycystic kidney disease; von Willebrands disease; forensic; h
tuberous sclerosis; hereditary hemorrhagica telangiectasia;
familial colonic polyposis; osteogenesis imperfecta; porphyria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Polymorphism; hypertension; agammaglobulinemia; diabetes insipidus;
Lesch-Nyhan syndrome; muscular dystrophy; Wiskott-Aldrich syndrome;
Fabrys disease; familial hypercholesterolemia; hereditary spherocytosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Polymorphic fragment of hypertension associated gene APOC4
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insipidus, Lesch-Nyhan syndrome, muscular uysuren, manyong syndrome, Fabrys disease, familial hypercholesterolemia, polycystic syndrome, Fabrys disease, familial hypercholesterolemia, polycystic kidney disease, hereditary spherocytosis, von Willebrands disease, tuberous sclerosis, hereditary hemorrhagica telangiectasia, familial
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03-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim
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nilarity 96.0%;
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99US-00304232.
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No. 9.7e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            the diagnosis
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                     The present invention describes a method for detecting the presence polymorphisms associated with inflammatory bowel diseases such as ulcerative colitis and Crohn's disease. The methods can be used to d the presence of genetic polymorphisms associated with inflammatory b disease and correlating their occurrence with disease states. They used in this way for phenotypic correlations, forensics, paternity testing, medicine and genetic analysis. The present sequence is a polymorphic site described in the exemplification of the invention
                                                                                                                                                  Testing for the prese
bowel disease, using
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; inflammatory bowel disease; Crohn's disease; ulcerative colitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human inflammatory bowel disease associated
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10-APR-2000; 2000US-0196046P
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   single nucleotide
                                                                                                                            Claim 1; Page 61; 463pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           posis, Ehlers-Danlos syndrome, osteogenesis imperfecta, ittent porphyria. The polymorphic forms can also be use identify individuals
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      polymorphism; SNP; chromosome 19p13; paternity test;
3; forensic test; gene therapy; ds.
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Pred. No. 9.
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                                                                                                                                                                                                               The present invention describes a method for detecting the presence polymorphisms associated with inflammatory bowel diseases such as ulcerative colitis and Crohn's disease. The methods can be used to othe presence of genetic polymorphisms associated with inflammatory k disease and correlating their occurrence with disease states. They used in this way for phenotypic correlations, forensics, paternity testing, medicine and genetic analysis. The present sequence is a polymorphic site described in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                           Testing for the bowel disease,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       single nucleotide polymorphism; SNP; chromosome 19p13; paternity test; chromosome 5q31-33; forensic test; gene therapy; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; inflammatory bowel disease; Crohn's disease; ulcerative colitis;
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                                              ADA74797
                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                       Claim 1;
                                                                                                                                                                                                                                                                                                                                                                             WPI; 2001-367874/38.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         10-DEC-1999; 99US-0170257P.
10-APR-2000; 2000US-0196046P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          11-DEC-2000; 2000WO-US033632
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14-JUN-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   misc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            09-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAH91536
                                                                                                                                                                                                                                                                                                                                                                                                                       (WHED )
                                                                                                                                                 Local Similarity
                                                                                                                          530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    inflammatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29
                                                                                                      29
                                                                                                                                                                                                                                                                                                                                           for the presence of polymorphisms associated with inflammatory isease, using a hybridization assay.
                                                                                                                                                                                                                                                                                                                                                                                                                       WHITEHEAD INST BIOMEDICAL ELLIPSIS BIOTHERAPEUTICS C
                                                                                                                                                                                              29
                                                                                                                                                                                                                                                                                                                     Page
                                              standard;
                                                                                                                          GCATCCTCCTGCCTCAGCCTCCCAAGTAG 558
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            B₽;
                                                                                                                                                 Conservative
  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                     64; 463pp; English.
                                                                                                                                                                                            6 A;
                                                                                                                                                                                                                                                                                                                                                                                                 ŗŢ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA;
                                              DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   bowel disease associated polymorphic site #611
                                                                                                                                                         2.3%;
                                                                                                                                                                                            5 C; 11
                                                                                                                                                                                                                                                                                                                                                                                                  Lander ES,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    a
"SNP, optionally T or C
                                              23
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                                              BP
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                                                                                                                                                Pred. No. le+
                                                                                                                                                          Score 23.2;
Pred. No. 1e
                                                                                                                                                                                           G; 6 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                  Rioux J,
                                                                                                                                                                                                                                                                                                                                                                                                                        CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                   RES.
                                                                                                                                                          1e+03;
                                                                                                                                                                      В
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                                                                                                                                                                    Length 29;
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                                                                                                                                               Gaps
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RESULT 339
AAH91561/c
HX SX XW XX DXX AX X II
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                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                    Human; inflammatory bowel disease; Crohn's disease; ulcerative cosingle nucleotide polymorphism; SNP; chromosome 19p13; paternity chromosome 5q31-33; forensic test; gene therapy; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                         for ethnic affiliation and thus for estimating ethnic affiliation, as well as for determining an ethnic specific haplotype, estimating genetic affiliation and during forensic analysis. The 3' A-rich region of the Alu repeat used within the method is a 'hot spot' for diversity making this region very useful for forensic analysis. The current sequence is that of the PCR primer F1209 of the invention which was used to sequence the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               achieved via obtaining haplotypes from specific DNA sequence analysis comparing haplotypes from human donors of known ethnic origin with expected haplotype frequency. The DC is used as an ethnic descent predictor. The method of the invention may be useful for determining a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Determining discriminant function coefficient for ethnic affiliation, comprises comparing haplotypes from donors of known ethnic origin with expected haplotype frequency and estimating coefficient from obtained
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to a method which comprises determining discriminant function coefficient (DC) for ethnic affiliation.
                                                                                                               Human inflammatory bowel disease associated polymorphic site #636.
                                                                                                                                                                                                       AAH91561 standard;
                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 2; Col 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                27-OCT-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            27-OCT-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          08-APR-2003.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          discriminant
                          Homo sapiens
                                                                                                                                              09-OCT-2001
                                                                                                                                                                          AAH91561;
                                                                                                                                                                                                                                                                                                                                                                                                                               human LDLR DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-566586/53.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Deininger P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PCR primer F1209 used to sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (DEIN/) DEININGER P.
(KASS/) KASS D.
                                                                                                                                                                                                                                                                                                            861 AGTGCTGGGATTACAGGCGTGAG 883
                                                                                                                                                                                                                                                                               23
                                                                                                                                                                                                                                                                                                                                         23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                c predictor;
low density
                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                               AGTGCTGGGATTACAGGCGTGAG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               density lipoprotein
                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                  BP; 5 A; 10 C; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             function coefficient; DC; ethnic affiliation; haplotype;
ictor; forensic analysis; Alu repeat; hot spot; diversity
                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               97US-00958009.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      25pp; English.
                                                                                                                                                                                                       DNA;
                                                                                                                                                                                                                                                                                                                                                       2.3%;
                                                                                                                                                                                                       24
                                                                                                                                                                                                       BP.
                                                                                                                                                                                                                                                                                                                                         <u>,,</u>
                                                                                                                                                                                                                                                                                                                                                       Score 23;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                  G; 5 T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               receptor; LDLR; ss; primer;
                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       human LDLR DNA
                                                                                                                                                                                                                                                                                                                                                       DB 1; Le . 8.9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                  U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                   Length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                diversity;
CR; F1209.
                                                                      colitis;
ty test;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This is
                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  a DC
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Кey

Location/Qualifiers

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RESULT 340
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Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 24 BP; 5 A; 4 C; 11 G; 3 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention describes a method for detecting the presence of polymorphisms associated with inflammatory bowel diseases such as ulcerative colitis and Crohn's disease. The methods can be used to detect the presence of genetic polymorphisms associated with inflammatory bowel disease and correlating their occurrence with disease states. They may be used in this way for phenotypic correlations, forensics, paternity testing, medicine and genetic analysis. The present sequence is a polymorphic site described in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Testing for the presence of polymorphisms associated with inflammatory bowel disease, using a hybridization assay.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Daly M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10-DEC-1999; 99US-0170257P.
10-APR-2000; 2000US-0196046P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       11-DEC-2000; 2000WO-US033632
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO200142511-A2
                                                                                                                                          EP450370-A.
                                                                                                                                                                                    Branch migration; linker; displacer; target;
                                                                                                                                                                                                           X-T-D oligonucleotide
                                                                                                                                                                                                                                   19-DEC-1991
                                                                                                                                                                                                                                                                              AAQ14732 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (WHED )
          WPI; 1991-297200/41
                                Wetmur
                                                                          26-MAR-1990;
                                                                                                15-MAR-1991;
                                                                                                                     09-OCT-1991.
                                                      (ENZO-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2001-367874/38.
                                                                                                                                                                                                                                                                                                                                                           839 TCTGCCTGCCTCGGCCTCCCAAAG 862
                                                                                                                                                                                                                                                                                                                                     24
                                                                                                                                                                                                                                                                                                                                                                               23;
                                JG,
                                                     ENZO
                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WHITEHEAD INST BIOMEDICAL RES. ELLIPSIS BIOTHERAPEUTICS CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Page 65; 463pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hudson TJ,
                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                               Quartin
                                                      BIOCHEM INC
                                                                                                                                                                                                                                  (first entry)
                                                                           90US-00499938
                                                                                                  91EP-00104066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        12
/*tag= a
/note= "SNP, optionally T or C at this position"
                                                                                                                                                                                                                                                                               DNA;
                                                                                                                                                                                                                                                                                                                                                                                          2.3%;
95.8%;
                                RS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Lander ES,
                                                                                                                                                                                                                                                                                 26
                               Engelhardt DL;
                                                                                                                                                                                                                                                                                 ₽₽
                                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                            Score 23;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Rioux J,
                                                                                                                                                                                                                                                                                                                                       \vdash
                                                                                                                                                                                                                                                                                                                                                                                            9.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                        DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                     Length 24;
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Matches Best Query Match

24;

Conservative

<u>.</u>

Mismatches

Indels

0,

Gaps

0

Local

Similarity

2.3%;

Score 22.8; DB 1; Pred. No. 9.8e+02;

DB 1;

Length 26,

Sequence

26

BP; 5

Α.

9 C; 8 G; 4 T; 0 U;

0 Other;

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RESULT 341
AAQ71189/c
ID AAQ711
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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local
                      To obtain vector p9lox5, a SalI cohesive-ended duplex containing a lox sequence from the phage P1 lox/Cre recombinase system was made by annealing oligos B16 (AAQ71187) and B17 (AAQ71188) and cloning into SalI-cleaved pUC9. Primer 614 (AAQ71189), designed from a consensus Alu repeat sequence, was used to prime amplification of the inter-Alu region of genomic DNA from line B2.13, which contains human chromosome-11 in a CHO background. The product was ligated to p9lox5 by turbo cloning. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The sequences represented in AAQ14004-32 and AAQ14728-33 are used in examples of the specification, illustrating the method of the inventi Oligonucleotides AAQ14016-29 and AAQ14728-33 are linkers, displacers targets. Nomenclature: X=XmaI restriction site T=target D=polarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Branch migration of nucleotide(s) - using one strand of recipient polydeoxy-nucleotide sequence and displacer sequence of single-stranded DNA
                                                                                                                                                                                                                                              Boyd
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 26 BP; 3 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page 15;
                                                                                                                                                                          Cloning of DNA molecules - by ligating with a vector contg cyclisation with Cre protein and dissociation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25-MAR-2003
29-MAR-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAQ71189;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAQ71189
                                                                                                                                                Disclosure; Page 7; 36pp; English.
                                                                                                                                                                                                                   WPI; 1994-279754/34.
                                                                                                                                                                                                                                                                                                                                11-FEB-1994;
                                                                                                                                                                                                                                                                                                                                                           18-AUG-1994.
                                                                                                                                                                                                                                                                                                                                                                                    WO9418333-A1
                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                      Vector; p9lox5; turbo cloning; lox; lox/Cre recombinase; pUC9;
inter-Alu region; polymerase chain reaction; PCR; primer; ampl
human chromosome-11; CHO; Chinese hamster ovary; 88.
                                                                                                                                                                                                                                                                                                     12-FEB-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Inter-Alu region primer 614.
                                                                                                                                                                                                                                                                          (MEDI-) MEDICAL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             standard;
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(first en
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                                                                                                                                                                                                                                                                           RES COUNCIL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    34pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               26
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Pred. No. 9.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             993
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          amplification;
                                                                                                                                                                                           a lox site,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  invention.
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     RESULT 342
26. MAR. 2001; 2001US-0278894P
27. MAR. 2001; 2001US-0279344P
28. MAR. 2001; 2001US-0279344P
28. MAR. 2001; 2001US-0279344P
30. MAR. 2001; 2001US-027934P
30. MAR. 2001; 2001US-0280233P
02. APR. 2001; 2001US-0280802P
02. APR. 2001; 2001US-0280802P
02. APR. 2001; 2001US-0280900P
04. APR. 2001; 2001US-0281194P
03. APR. 2001; 2001US-028166P
03. APR. 2001; 2001US-0288742P
03. MAY. 2001; 2001US-0288342P
03. MAY. 2001; 2001US-028129P
16. MAY. 2001; 2001US-028149P
16. MAY. 2001; 2001US-029129P
16. MAY. 2001; 2001US-029129P
16. MAY. 2001; 2001US-029139P
16. MAY. 2001; 2001US-029139P
16. MAY. 2001; 2001US-029139P
16. MAY. 2001; 2001US-029139P
16. JUN. 2001; 2001US-029139P
18. JUN. 2001; 2001US-0299027P
19. JUN. 2001; 2001US-0299303P
19. JUN. 2001; 2001US-0299303P
16. AUG. 2001; 2001US-0299303P
16. AUG. 2001; 2001US-0399319P
31. JUN. 2001; 2001US-0399319P
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20-MAR-2001;
21-MAR-2001;
22-MAR-2001;
23-MAR-2001;
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08-MAR-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human NOV-associated probe from primer-probe set Ag3293
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20-MAY-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    human; probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NOVX; cytostatic; cardiant; antiarteriosclerotic; antiasthmatic; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                20-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 hypotensive; cardiomyopathy; bronchial asthma; gene therapy; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             16-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  L3-MAR-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    standard; DNA;
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2001US-0275601P.
2001US-027600P.
2001US-0276776P.
2001US-027732P.
2001US-027732P.
2001US-027733P.
2001US-027733P.
2001US-027733P.
2001US-027733P.
2001US-027733P.
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2001US-0274194P.
2001US-0274281P.
2001US-0274322P.
2001US-0274323P.
2001US-0275235P.
2001US-0275238P.
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Human NOV20a

RTQ-PCR probe.

(first

entry)

Human; ss; PCR; NOVX; diabetes; obesity; infectious disease; anorexia; cancer-associated cachexia; cancer; neurodegenerative disorder; Alzheimer's disease; Parkinson's disease; immune disorder;

RESULT 343

ADN62195

standard;

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ADN62195; ADN62195 밁

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ATGCACCACCACTCCTGGCTAATTTT

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Best Local S
Matches 24
                                                                                                                          This invention describes novel human NOVX polypeptides which have cytostatic, cardiant, antiarteriosclerotic, antiasthmatic and hypotensive activity. Pharmaceutical compositions comprising the NOVX proteins or nucleic acid molecules or NOVX antibodies are useful for preventing or treating a disorder associated with aberrant NOVX expression or activity e.g. cancer, hypertension, atherosclerosis, cardiomyopathy or bronchial asthma. The products of the invention can be used for gene therapy or in a vaccine. ABX13460-ABX13462 and ABX97186-ABX97933 represent PCR primers and probes used in the amplification and isolation of the NOVX polymorphic represented in ABX97080-ABX97185 which encode the polypeptides represented in ABX95041-ABU65218. The probes described in the invention are modified at the 5'-ard by more
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            14-NOV-2001;
14-NOV-2001;
14-NOV-2001;
14-NOV-2001;
21-NOV-2001;
03-DEC-2001;
03-DEC-2001;
04-DEC-2001;
03-JAN-2002;
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12-SEP-2001;
27-SEP-2001;
27-SEP-2001;
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31-OCT-2001;
14-NOV-2001;
                                                                                                                                                                                                                                                                                                                                                                     NOVX polypeptides and polynucleotides, useful for preventing or treating a disorder associated with aberrant NOVX expression or activity e.g., cancer, hypertension, atherosclerosis, cardiomyopathy or bronchial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Spaderna SK, Catterton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Padigaru M,
Zerhusen BD,
                                                                                                   Sequence 26 BP; 6 A; 9 C; 3 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                             Example C;
                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-723332/78.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Patturajan M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Lepley DM,
573 ATGCACCACTACACCTGGCTAATTTT 598
                                           24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURAGEN CORP.
                                                       Similarity
                                                                                                                                                                                                                                                                                                                             Page 621; 1103pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2001US-0318462P
2001US-0318770P
2001US-0325430P
2001US-0325681P
2001US-0335301P
2001US-0335301P
2001US-03352172P
2001US-0332172P
2001US-0332771P
2001US-0332771P
2001US-0332772P
2001US-033274P
2001US-033274P
2001US-0337426P
2001US-0337185P
2001US-0337185P
2001US-0337185P
                                                                                                                                                                                                                                                                                                                                                                                                                                                Spytek KA, Shenoy ov, Gusev V, Ji W, Gorman L, Miller Cb, Chernev V, Gangolli E, Vernet CAM, Guo X, Tchernev V, R, Casman SJ, Malyankar UM, Gerlach V, Liu Y, Catterton E, Burgess C, Leite M, Zhong H, /
                                         2.3%;
ilarity 92.3%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2002US-00092900
                                         0
                                                       Score 22.8;
Pred. No. 9
                                            Mismatches
                                                         .8e+02
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                                                                     Length
                                            Indels
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Alsobrook JP;
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                                         Gaps
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08-MAR-2001
09-MAR-2001
113-MAR-2001
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115-MAR-2001
120-MAR-2001
20-MAR-2001
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21-MAR-2001
22-MAR-2001
23-MAR-2001
27-MAR-2001
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21-MAY-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            07-MAR-2002; 2002US-00092900
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    2001US-0277338P.
2001US-0277318P.
2001US-027781SP.
2001US-02788194P.
2001US-027893P.
2001US-027993P.
2001US-0280822P.
2001US-0280822P.
2001US-0280822P.
2001US-0280822P.
2001US-02887424P.
2001US-02887424P.
2001US-02887424P.
2001US-02887424P.
2001US-02887424P.
2001US-02887424P.
2001US-02887424P.
2001US-02887424P.
2001US-02988728P.
2001US-0291290P.
2001US-0291290P.
2001US-029130P.
2001US-0294899P.
2001US-029489P.
2001US-029489P.
2001US-029489P.
2001US-029489P.
2001US-039310P.
2001US-0318742P.
2001US-0318462P.
2001US-0318462P.
2001US-0318462P.
2001US-03312903P.
2001US-0335301P.
2001US-0335301P.
2001US-0335301P.
2001US-0335301P.
2001US-0335301P.
2001US-0335301P.
2001US-0335301P.
2001US-0335301P.
2001US-0333184P.
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2001US-0276994P.
2001US-0277239P.
2001US-0277321P.
2001US-0277327P.
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2001US-0274849P.
2001US-027523BP.
2001US-0275578P.
2001US-0275579P.
2001US-0275609P.
2001US-0275600P.
The invention relates to an isolated polypeptide (designated NOVX, or CC NOV1-NOV127) comprising a sequence selected from 178 fully defined amino CC acid sequences (and their mature forms, variants and fragments). Also CC included are an isolated nucleic acid molecule encoding NOVX, a vector CC comprising the nucleic acid, a cell comprising the vector, methods for determining the presence or amount of the polypeptide or the nucleic acid molecule in a sample, methods for determining the presence of or predisposition to a disease associated with altered levels of expression of the above polypeptide or nucleic acid molecule in a first mammalian cc subject, a method for identifying an agent that binds to the above polypeptide, a method for identifying a potential therapeutic agent for use in the treatment of a pathology that is related to aberrant cc expression or physiological interactions of the polypeptide, a method of screening for a modulator of activity or of latency or predisposition to greening for a modulator of activity or of latency or predisposition to a pathology associated with the polypeptide and a method for modulating ct the activity of the polypeptide citted above. The composition and methods are useful for diagnosing, preventing or treating diseases such as cc ancer, neurodegenerative disorders like Alzheimer's disease or Parkinson's disease, immune disorders, haematopoietic disorders, cancer also used in Cc chromosome mapping, tissue typing, preventive medicine and cc pharmacogenomics. The polypeptides are also useful as vaccines. The compositione and cc pharmacogenomics is an RTQ-PCR (real time quantitative PCR) probe used to
             Padigaru M, S;
Zerhusen BD,
Patturajan M,
Fernandes ER,
Anderson DW,
Alsobrook JP,
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(LEIT/)
(ZHON/)
(ALSO/)
(ALSO/)
(LEPL/)
(RIEG/)
(BURG/)
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(GUSE/
(JIWW/
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(SHEN/)
(TAUP/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New NOVX polypeptides and nucleic acid molecules useful for diagnosing, preventing or treating NOVX-associated disorders, e.g. cancer, diabetes, infection or obesity, and in chromosome mapping, tissue typing or
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(LIUY/
(ANDE/
(SPAD/
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(PATT/
(GANG/
(VERN/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example C;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 pharmacogenomics
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MALYANKAR U M.
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SPADERNA S K.
CATTERTON E.
LEITE M W.
ZHONG H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GORMAN L.
MILLER C E.
KEKUDA R.
PATTURAJAN M.
GANGOLLI E A.
VERNET C A M.
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PENA C E A.
LI L.
ZERHUSEN B I
GUSEV V Y.
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SPYTEK K A.
SHENOY S G.
TAUPIER R J.
PENA C E A.
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RIEGER D K
BURGESS C
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LEPLEY D M.
RIEGER D K.
BURGESS C E.
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JI W.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Spytek KA, Shenoy SG, Taupie;
), Gusev VY, Ji W, Gorman L, I
M, Gangolli EA, Vernet CAM, Ge
IR, Casman SJ, Malyankar UM, Ge
I, Spaderna SK, Catterton E, Le
IP, Lepley DM, Rieger DK, Burg
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Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide CC primer extension (SNPE) primers, and the sequences of regions flanking cc sites of single nucleotide polymorphisms SNPs. The present invention cc includes kits for determining the presence or absence of a SNP, using the cligonucleotides of the invention. The PCR primers are used to amplify a SNP flanking sequence, the SNPE primer is used as genotyping primer. CC The oligonucleotides are useful for genotyping a nucleic acid sample by performing a single-nucleotide primer extension reaction. The collgonucleotides are useful for determining the presence, absence or clentity of a SNP and for genotyping nucleic acid sample by performing a single-nucleotide primer extension reaction. The cc identity of a SNP and for genotyping nucleic acid samples, for e.g. to assess by association analysis the genotype of an individual or group of individuals, having a pathological phenotypic trait suspected of being caused by one or more SNPs. Phenotypic traits include diseases e.g. cc agammaglobulinaemia, diabetes insipidus, Lesch-Nyhan syndrome, muscular dystrophy, familial hypercholesterolaemia, polycystic kidney disease, cc osteogenesis imperfecta and acute intermittent porphyria. Phenotypic traits also include symptoms of or susceptibility to multifactorial disease of which a component is or may be genetic such as autoimmune cd disease of which a component is or may be genetic such as autoimmune inflammation, cancer, nervous system diseases and infection by pathogenic microorganism. The method is also useful in forensic investigations and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local S
Matches 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genotyping; agammaglobulinaemia; diabetes insinidns. carrar.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              assay tissue specific expression of a NOVX mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SNP specific
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 1; Page 54; 83pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15-OCT-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        inflammation;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            stic kidney disease; osteogenesis imperfecta; autoimmune disease; intermittent porphyria; rheumatoid arthritis; multiple sclerosis;
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Pred. No. 9.
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cc includes kits for determining the presence or absence of a SNP, using the colligonucleotides of the invention. The PCR primers are used to amplify a CSNP flanking sequence, the SNPE primer is used as a genotyping primer. CC The oligonucleotides are useful for genotyping a nucleic acid sample by colligonucleotides are useful for genotyping a nucleic acid sample by colligonucleotides are useful for determining the presence, absence or colligonucleotides are useful for determining the presence, absence or clantity of a SNP and for genotyping nucleic acid samples, for e.g. to cassess by association analysis the genotype of an individual or group of individuals, having a pathological phenotypic trait suspected of being caused by one or more SNPs. Phenotypic traits include diseases e.g. cc agammaglobulinaemia, diabetes insipidus, Lesch-Nyhan syndrome, muscular consteogenesis imperfecta and acute intermittent porphyria. Phenotypic traits also include symptoms of or susceptibility to multifactorial consteogenesis include acomponent is or may be genetic such as autoimmune cc disease of which a component is or may be genetic such as autoimmune cc diseases, including, rheumatoid arthritis, multiple sclerosis,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      English.
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Pred. No. 1
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Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide CC primer extension (SNPE) primers, and the sequences of regions flanking CC sites of single nucleotide polymorphisms SNPs. The present invention CC includes kits for determining the presence or absence of a SNP, using the CC oligonucleotides of the invention. The PCR primers are used to amplify a CC sNP flanking sequence, the SNPE primer is used as a genotyping primer. CC the oligonucleotides are useful for genotyping a nucleic acid sample by CC performing a single-nucleotide primer extension reaction. The CC oligonucleotides are useful for determining the presence, absence or CC identity of a SNP and for genotyping nucleic acid samples, for e.g. to assess by association analysis the genotype of an individual or group of CC individuals, having a pathological phenotypic trait suspected of being CC caused by one or more SNPs. Phenotypic traits include disease e.g. CC osteogenesis imperfecta and acute intermittent porphyria. Phenotypic ctraits also include symptoms of or susceptibility to multifactorial
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       س
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
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RESULT 347
AAH91322/c
ID AAH91
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          disease of which a component is or may be genetic such as autoimmune diseases, including, rheumatoid arthritis, multiple sclerosis, inflammation, cancer, nervous system diseases and infection by pathogeni microorganism. The method is also useful in forensic investigations and paternity analysis. The present sequence represents a single nucleotide primer extension (SNPE) primer specific for a human SNP containing DNA
                     The present invention describes a method for detecting the presence polymorphisms associated with inflammatory bowel diseases such as ulcerative colitis and Crohn's disease. The methods can be used to d the presence of genetic polymorphisms associated with inflammatory b disease and correlating their occurrence with disease states. They used in this way for phenotypic correlations, forensics, paternity testing, medicine and genetic analysis. The present sequence is a polymorphic site described in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; inflammatory bowel disease; Crohn's disease; ulcerative cosingle nucleotide polymorphism; SNP; chromosome 19p13; paternity chromosome 5q31-33; forensic test; gene therapy; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAH91322 standard; DNA;
                                                                                                                                                                                                                                                                                         10-DEC-1999; 99US-0170257P
10-APR-2000; 2000US-0196046P
                                                                                                                                                                                                                                                                                                                                                                                                                               misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        09-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAH91322;
                                                                                                                                                            Testing for th
bowel disease,
                                                                                                                                                                                                                          Daly M,
                                                                                                                                                                                                                                                                                                                              11-DEC-2000; 2000WO-US033632
                                                                                                                                                                                                                                                                                                                                                                                WO200142511-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human
                                                                                                                                   Claim 1; Page 55; 463pp; English
                                                                                                                                                                                                 WPI; 2001-367874/38
                                                                                                                                                                                                                                                    (BLLI-)
                                                                                                                                                                                                                                                                ( MHED )
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
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                                                                                                                                                                           for the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AGTGCTGGGATTACAGGCGTGAGCCAC 887
                                                                                                                                                                                                                           Hudson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AGTGCTGAAATTACAGNCGTGAGCCAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP; 8 A; 6 C; 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                            ne presence
using a hy
                                                                                                                                                                                                                                                                                                        99US-0170257P
                                                                                                                                                                                                                          IJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                         /*tag=
/note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 bowel disease
                                                                                                                                                            ence of polymorphisms
a hybridization assay
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      88.9%;
                                                                                                                                                                                                                             Lander ES,
                                                                                                                                                                                                                                                                                                                                                                                              a "SNP,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                       CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  associated
                                                                                                                                                                                                                             Rioux J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             22.8;
10, 1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0 U; 1 Other;
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                                                                                                                                                                            associated with inflammatory
                                                                                                                                                                                                                              Siminovitch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  polymorphic site
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Query Match

Sequence

27

B₽;

6 A;

4 C; 11 G;

5 T; 0 U; 1 Other;

Score

22.8;

DB 1;

Length

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RESULT 349
AA169885/c
ID AA1698
XX
AC AA1698
XX
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AAF92843
ID AAF928
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                                                                                                                                                Query Match
Best Local S
Matches 23
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Matches 24; Conserv
                                                                                                                                                                                                                  The present invention relates to a method for treating a patient diagnosed as having a lower than normal high density lipoprotein-cholesterol (HDL-C) level, a higher than normal triglyceride level, or a cardiovascular disease, involving administering a compound that modulates LXR- or RXR-mediated transcriptional activity or ABC1 expression or activity. The LXR gene product may be used in an assay to identify compounds useful for the treatment of a disease or condition selected a lower than normal HDL cholesterol level, a higher than normal triglyceride level, and a cardiovascular disease
              AAI69885;
                                                                                                                                                                                                                                                                                                                                              Disclosure; Fig 3; 317pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-244356/25.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  High density
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAF92843 standard; DNA; 24
                                    AAI69885 standard;
                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                   disease,
                                                                                                                                                                                                                                                                                                                                                                                                        Treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hayden MR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-SEP-2000; 2000WO-IB001492.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200115676-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human ABC1 transcription factor binding site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17-MAY-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAF92843;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (XENO-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (UYBR-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1003
                                                                                                                       208
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XENON GENETICS INC.
                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                                      by administering a compound that modulates LXR-ptional activity.
                                                                                                                                                                                                                                                                                                                                                                                              a lower than normal high density lipoprotein-cholesterol (HDL-C) higher than normal triglyceride level, or a cardiovascular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AGCGATTCTCCTGTCTCAGCCTCCCAA 1029
                                                                                                                       AGGCTGGTCTCGAACTCCCGACCT
                                                                                                                                                                                              24 BP; 4 A; 8 C; 6 G; 6 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AGCGATTCTTCNGCCTCAGCCTCCCAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Brooks-Wilson AR,
                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2000US-00526193.
2000US-0213958P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    lipoprotein-cholesterol; HDL-C; cardiovascular;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            99US-0151977P.
                                    DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         88.9%;
                                                                                                                                                         95.8%;
                                    24
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                                    ВP
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                                                                                                                                            Pred. No. 9.76
0; Mismatches
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0; Mismatches
                                                                                                                                                            Score 22.4;
Pred. No. 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pimstone
                                                                                                                                                                                                                                                                                                                                                                                                                                                          SN,
                                                                                                                                                            .7e+02;
                                                                                                                                                                                                0 Other;
                                                                                                                                                                        DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                             or a cardiovascular
                                                                                                                                                                     Length 24;
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                                                                                                                                                                                                                                                                                                                                                                                   or RXR-mediated
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABC1;
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                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                             RESULT 350
                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention relates to human transglutaminase 12 (see AAG78882). The transglutaminase and its coding sequence are useful in the diagnosis and treatment of malignant tumours, haemopathy, HIV infection, immunological diseases and various inflammations. The present sequence is a PCR primer which was used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human transglutaminase 12 and encoded polynucleotide, used in diagnoused treatment of malignant tumors, hemopathy, human immunodeficiency virus infection, immunological diseases and inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; transglutaminase 12; cytostatic; virucidal; immunomodulatory; antiinflammatory; haemostatic; gene therapy; malignant tumour; haemopathy; HIV infection; immunological disease; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           14-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 24 BP; 5 A; 8 C; 4 G; 7 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200170787-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      haemopathy; HIV PCR primer; ss.
                                                                                                                                                                     Human; RAD51;
                                                                                                                                                                                              PCR primer
                                                                                                                                                                                                                                              AAS00333;
                                                                                                                                                                                                                                                                     AAS00333 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 2; Page 17; 37pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-611474/70.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Мао У,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      26-FEB-2001; 2001WO-CN000243
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
Wang WW,
                                                                       08-SEP-2000;
                                                                                                15-MAR-2001
                                                                                                                                                                                                                       17-MAY-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                10-MAR-2000; 2000CN-00111967.
                        (USSH ) US DEPT HEALTH & HUMAN SERVICES
                                                10-SEP-1999;
                                                                                                                       WO200118254-A2
                                                                                                                                                                                                                                                                                                                                              388 CAAAGTGCTGGGATTACAGGCGTG 411
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   transglutaminase 12 PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       infection, immunological diseases and
                                                                                                                                                                                                                                                                                                                                  24
                                                                                                                                                                                                                                                                                                                                                                                 23;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SHANGHAI BIOWINDOW GENE DEV
                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                              #2,
                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                        2000WO-US024786
                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                        breast
                                                                                                                                                                                               used to amplify human RAD51
                                               99US-0153288P
                                                                                                                                                                                                                                                                      DNA;
                                                                                                                                                                                                                                                                                                                                                                                              2.3%;
95.8%;
                                                                                                                                                                        cancer;
                                                                                                                                                                                                                                                                        24
                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                        BRCA1; BRCA2; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                              Score 22.4;
Pred. No. 9.
                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   #1
                                                                                                                                                                                                                                                                                                                                                                                             9.7e+02
                                                                                                                                                                                                                                                                                                                                                                                                            DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                  0 Other;
                                                                                                                                                                                              gene at position -2339.
                                                                                                                                                                                                                                                                                                                                                                                                          Length 24;
                                                                                                                                                                                                                                                                                                                                                                                   Indels
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Gaps

WPI; 2001-235217/24

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The sequence represents PCR primer #2, used to amplify human RAD51 gene at position -2339 upstream of the transcription start site of human RAD51 gene. The RAD51 gene is useful in diagnosing genetic predisposition or susceptibility to breast cancer in an individual using the following steps: (1) detecting a mutation in the RAD51 gene in a human subject, comprising analysing a sample from the subject to detect the mutation; (2) assessing the risk of developing breast cancer, comprising; (a) analysing a sample from the subject for the presence of BRCA1 and/or BRCA2 mutations; and (b) if (a) is positive, analysing the sample for a mutation in the RAD51 gene, where the presence of the RAD51 mutation in compared to a subject having at least one of the BRCA mutations and a wild-type RAD51 gene. Primers derived from the sequence can be used in a kit for detecting a mutation in the RAD51 gene of a subject, which is associated with a predisposition to breast cancer, comprising at least 2 nucleic acid primers derived from the RAD51 gene sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New nucleic acids comprising a mutant of the RAD51 gene, useful for diagnosing genetic predisposition or susceptibility to breast cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 24 BP; 5 A; 10 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 42; 42pp; English.
                                                                                                                                                                                                                                                                                                                                                                                     Zinc finger haemopathy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABZ21100 standard; DNA;
                                                                                                                                                                                                                              06-NOV-2000; 2000CN-00127270
                                                                                                                                                                                                                                                             06-NOV-2000; 2000CN-00127270
                                                                                                                                                                                                                                                                                            05-JUN-2002.
                                                                                                                                                                                                                                                                                                                                                         Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                     Zinc finger
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABZ21100
                                                                                        New zinc finger protein 54.67 polypeptide for treating malignant tumors, inflammations, immunological diseases, hemopathy and human
                                                                                                                                       WPI; 2002-699446/76
                                                                                                                                                                                                 (BODE-) BODE GENE DEV CO LTD
                                                                           immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        538
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       23;
                                                                                                                                                                    Xie
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CTGCCTCAGCCTCCCAAGTAACTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CTGCCTCAGCCTCCCAAGTAGCTG 561
                                                                                                                                                                                                                                                                                                                                                                                      protein 54.67; tumour; inflammation;
HIV infection; cytostatic; anti-HIV;
                                                                                                                                                                                                                                                                                                                                                                                                                                   protein 54.67 PCR primer #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2.3%;
95.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ٥,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 22.4;
Pred. No. 9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1; Length 24;
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Indels

<u>;</u>

0

immunological disease; PCR; primer; ss.

Example 2;

Page

16

(Disclosure); 34pp; Chinese.

present invention relates to zinc finger protein 54.67 (ABB98889) zinc finger protein can be used for treating various diseases, su

such

Sequence 24

BP; 3 A; 9 C;

7 G;

5 T;

0 U;

0 Other;

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RESULT 352
ABA96912/c
ID ABA969
XX ABA969
XX ABA969
XX ABA969
XX ABA969
XX Homan
DT 15-MAY
XX Human
XX Human
XX Human
XX Immunx
XX Immunx
XX Immunx
XX Homo i
XX I Human
PT 1-MA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    뭉
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                                              The invention relates to human argininase 9 (AAM49102), nucleic acids commoding it (ABA96911), and a method for the recombinant production of CC argininase 9. The protein has a molecular weight of 9 kD and has homology cover a 60 amino acid stretch to the protein fragment shown in AAM49105. CC The present invention additionally discloses an antagonist of argininase CC argininase 9. Argininase 9, and an antibody which specifically binds to argininase 9. Argininase 9, and nucleotides which encode it may be used CC argininase or urea metabolism, developmental disorders, malignant tumours, compared to screen for modulators of its activity or for peptide CC used to screen for modulators of its activity or for peptide CC immune disorders and inflammatory conditions. The protein may also be CC used to screen for modulators of its activity or for peptide CC for nucleic acid amplification reactions or as a probe for hybridisation CC exemplification, or in producing gene chips or microarrays. Sequences ABA96912-CC ABA96913 represent reverse transcription-PCR (RT-PCR) primers used in an CC exemplification of the invention to isolate human argininase 9 cDNA. CC ID NO:3 (ABA96914) which is given on page 12 of the specification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 24 BP; 4 A; 7 C; 8 G; 5 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human argininase 9 and encoding polynucleotide, used in diagnosis and treatment of malignant tumors, hemopathy, human immunodeficiency virus infection, immunological diseases and inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            arginine metabolism disorder; urea metabolism disorder; developmental disorder; malignant tumour; cancer; gene therapy; immune disorder; inflammatory condition; cytostatic; antiinflam
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        27-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; argininase 9; recombinant production; argininaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABA96912 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2002-090440/12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              мао Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19-MAY-2000; 2000CN-00115753.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14-MAY-2001; 2001WO-CN000788
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human argininase 9 RT-PCR primer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 3; Page 29; 33pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (BIOW-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 lmmunomodulator; reverse transcription-PCR; RT-PCR; primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     721 GCCTCCTGAGTAGCTGGGACTACA 744
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Xie Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BIOWINDOW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GCCTCCCGAGTAGCTGGGACTACA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENE DEV INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2.3%;
95.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 22.4;
Pred. No. 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SHANGHAI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQ ID NO:3 version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         9.7e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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RESULT 354
ABT08420
ID ABT084
XX
XX
AC ABT084
XX
DT 27-NOV
XX
DE Human
XX
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ABQ83933/c
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Best Local S
Matches 23
                                                                                                                                                                                   Query Match
Best Local S
Matches 23
                                                                                                                                                                                                                                                  The present invention describes human breast susceptible gene coded protein 10.45 (I). Also described is a process for preparing (I) using DNA recombination techniques. (I) can be used for treating several diseases e.g. embryotic development deformity and tumours. The present sequence represents a PCR primer for (I), which is used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                          A novel human breast susceptible gene coded protein 10.45 polypeptide, and the polynucleotide encoding it, useful for treating several diseases e.g. embryotic development deformity and tumors.
                                                                                                                                                                                                                                                                                                                                                                                                                               Mao
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          embryotic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABQ83933 standard; DNA; 24 BP
           Human PSF promoter PCR primer
                                                                                                                                                                                                                                Sequence 24 BP; 9 A; 7 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-529778/57.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                12-SEP-2000; 2000CN-00125173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       03-APR-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; breast
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  04-FEB-2003
                                  27-NOV-2002
                                                                               ABT08420 standard;
                                                                                                                                                                                                                                                                                                                                   Example 2; Page 18 (Disclosure); 34pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      (BODE-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           868
                                                                                                                                                             192
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           breast
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      24
                                                                                                                                        24
                                                                                                                                                                                    23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    BODE GENE DEV CO LTD SHANGHAI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                              Similarity
                                                                                                                                                            TTTCTCCATGTTGGTCAGGCTGGT 215
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GGATTACAGGCGTGAGCCACCGCG 1
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                                                                                                                                                                                    Conservative
                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2000CN-00125173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          susceptible gene protein 10.45 PCR primer 2 SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      susceptible gene coded
elopment deformity; PCR
                                                                              DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2.3%;
                                                                                                                                                                                             2.3%;
                                                                               24
                                                                               ВP
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                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 22.4; DB 1;
Pred. No. 9.7e+02;
0; Mismatches 1;
                                                                                                                                                                                             Score 22.4; DB 1;
Pred. No. 9.7e+02;
             SEQ
                                                                                                                                                                                     Mismatches
             IJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         protein 10.45; tumour; primer; ss.
             NO:
             55
                                                                                                                                                                                                         DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length
                                                                                                                                                                                                        Length 24;
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                                                                                                                                                                                     Indels
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X F X P X P X S X X X X D X P X A X X

primer;

18-DEC-2002; 2002WO-US040698

Homo sapiens. WO2003054160-A2 03-JUL-2003. Human; alphoid;

immunodeficiency virus; HIV; anti-HIV;

latency;

genomic

DNA

primer Alu.

06-NOV-2003

(first entry)

ACF05122;

H

ACF05122 standard; DNA;

24

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RESULT 355
ACF05122/c
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                                                                                                                                                                                                                                                                      Query Match
Best Local S
Matches 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention relates to a recombinant expression construct encoding a reporter gene operably linked to a promoter from a mammalian gene induced by a cyclin-dependent kinase (CDK) inhibitor. The construct is useful for identifying compounds that inhibit the induction of genes induced by CDK inhibitors. The compounds are useful for preventing or treating a disease caused by CDK inhibitor induced gene expression, e.g. cancer other than colon cancer, renal failure, Alzheimer's disease, amyloidosis, age-related diseases, atherosclerosis or arthritis. The present sequence is a PCR primer used to amplify a human promoter suitable for use in the construct of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; cyclin-dependent kinase; CDK; cyclin-dependent kinase inhibitor; inhibitor; cancer; age-related disease; promoter; atherosclerosis; cytostatic; antiarteriosclerotic; nootropic; neuroprotective; nephrotropic; antiarthritic; arthritis; renal disease; Alzheimer's disease; amyloidosis; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-FEB-2001; 2001US-0265840P
21-MAY-2001; 2001US-00861925
                                                                                                                                                                                                                                                                                                                                                                                                Sequence 24 BP; 7 A; 2 C; 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 8; Page 130; 137pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          that inhibit the induction inhibitors for preventing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New recombinant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Poole
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-FEB-2002; 2002WO-US002784.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200266681-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (UNII)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2002-674960/72.
                                                                                                                                                                                   859 AAAGTGCTGGGATTACAGGCGTGA
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                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Roninson IB,
                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ant expression construct, useful for identifying conthe induction of genes induced by cyclin-dependent or preventing or treating cancer, renal failure or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ILLINOIS FOUND.
                                                                                                                                                                                                                                                                                                      2.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chang
                                                                                                                                                                                                                                                                      <u>.</u>
                                                                                                                                                                                                                                                                                                      Score 22.4; DB 1;
Pred. No. 9.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                          G; 5 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               8
                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                          24
                                                                                                                                                                                                        882
                                                                                                                                                                                                                                                                                                                                                                                                             0 Other;
                                                                                                                                                                                                                                                                                                                                            ۲.
                                                                                                                                                                                                                                                                                                                                        Length
                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                            24;
                                                                                                                                                                                                                                                                          0,
                                                                                                                                                                                                                                                                             Gaps
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ACF35685/c
ID ACF356
XX ACF3566
XX ACF356
XX ACF366
XX A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       á
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local S
Matches 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present sequence is that of primer Alu (EV1255) for human genomic DNA. This primer was used with primer A (see ACCO5121) in alphoid PCR amplifications that demonstrated preferential HIV integration in or near alphoid DNA in latently infected Jurkat cells. The invention provides isolated cells that harbour a latent immunodeficiency virus that is transcription competent, that can be reactivated, and that is an in vitro model for latent HIV infection in vivo. The cells are useful for investigating the nature of latent HIV. Such agents are useful for reducing the reservoir of latent HIV. Such agents are useful for reducing the reservoir of latent HIV. Methods are provided of treating an immunodeficiency virus infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel isolated cells that comprise transcription competent immunodeficiency virus e.g. HIV-1, or immunodeficiency virus-based retroviral vector integrated into its genome, useful for identifyir latent HIV activators.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Verdin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 19-DEC-2001; 2001US-0341727P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 24 BP; 4 A; 7 C; 9 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Trans-Golgi network integral membrane protein; cytostatic; antiinflammatory; immunomodulator;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 1; Page 33; 71pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (REGC ) UNIV CALIFORNIA
                                                            Detecting apoptosis in a cell, useful for treating cancer, an inflammatory disease, an autoimmune disease or a neurodegenerative disease, comprises detecting a decrease in TGNP activity or expres
                                                                                                                                                                                                                                                                                                                                                 13-DEC-2001; 2001GB-00029846
                                                                                                                                                                                                                                                                                                                                                                                                           13-DEC-2002; 2002WO-GB005670
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO2003050302-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          nootropic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human TGNP promoter amplifying forward primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ACF35685 standard;
                                                                                                                                                                              WPI; 2003-532920/50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2003-577369/54.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    541 CCTCAGCCTCCCAAGTAGCTGGGA 564
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                                                                                                                                                                                                                                                                                              EIRX THERAPEUTICS LTD
                                                                                                                                                                                                                                        Cotter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Jordan A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          gene therapy; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                  ŗ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2.3%;
95.8%;
                                                                                                                                                                                                                                        Murphy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 22.4;
Pred. No. 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          primer;
                                                                                                                                                                                                                                        Seery
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    9.7e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           neuroprotective;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TGNP; chromosome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           identifying
                                                               or expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           human;
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The invention relates to detecting apoptosis in a cell. The method involves detecting a decrease in trans-Golgi network integral membrane protein (TGNP) activity or expression by detecting the decrease in TGNP polypeptide or its homologue, a nucleic acid encoding the polypeptide, a nucleic acid that hybridizes under stringent conditions to the aforementioned nucleic acid, or their complements. The method, polypeptides, nucleic acid, and modulators are useful for treating cancer, an inflammatory disease, an autoimmune disease or a neurodegenerative disease. The present sequence represents a PCR primer for amplifying the human TGNP promoter
Sequence 24 BP; 6 A; 6 C; 8 G; 4 T; 0 U;
                     0 Other;
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XSSSSSSSSSSXXX

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Query Match
Best Local S
Matches 23
635 CTCTGTCACCCAGGCTGGAGTGCA 658
              23;
                       Similarity
                Conservative
                       2.3%;
                0
                       Score
Pred.
                 Mismatches
                         22.4;
No. 9
                        DB 1;
).7e+02;
                               Length
                  Indels
                                24;
                  0
                 Gaps
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0;

맑 S

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26-FEB-2004 (first entry)
PCR primer
                                                                         ADG28972 standard;
                                                                                                                                         24
SEQ ID 55 used to amplify human PSF promoter
                                                                            DNA;
                                                                            24
                                                                            ВP
   DNA.
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virucide; cytostatic; neuroprotective; nootropic; antiarteriosclerotic; antiarthritic; nephrotropic; viral infection; cancer; renal; age-related disease; Alzheimer's; atherosclerosis; arthritis; recombinant expression construct; cyclin-dependent kinase inhibitor;

gene therapy; human; Alzheimer's; atherosclerosis; arthritis; ss; PCR; primer; PSF promoter.

Homo sapiens

04-SEP-2003:

29-AUG-2002; 2002WO-US027584

29-AUG-2001; 2001US-0315791P

VIND ILLINOIS FOUND.

Roninson IB, Poole J;

WPI; 2003-731624/69.

New recombinant expression construct for identifying and modulating expression of genes regulated by cyclin-dependent kinase inhibitors, as genes involved in viral infection, cancer, renal diseases or agesuch

related diseases.

Example 8; SEQ ID NO 55; 143pp; English.

RESULT 357
ADG28972
IID ADG288
XX ADG288
XX ADG288
XX ADG288
XX YECOMI
XX ADG288
XX YECOMI
XX WPI;
XX WEW YECOMI
XX NEW YECOMI
X The invention relates to a novel recombinant expression construct encoding a reporter gene operably linked to a promoter from a manma viral or cellular gene induced by a cyclin-dependent kinase (CDK) inhibitor. The construct of the invention demonstrates virucide, cytostatic, neuroprotective, nootropic, antiarteriosclerotic, antiarthritic and nephrotropic activities and may be useful in identifying compounds that inhibit the induction of genes involved viral infection, cancer, renal diseases or age-related diseases incolved alzheimer's disease, atherosclerosis or arthritis, such genes being induced by cyclin-dependent kinase inhibitors. Furthermore, The conmany have gene therapy applications. The current sequence is that of being mammalian including

Example 11;

Page 80; 110pp; English

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SXS
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                                          This invention describes a novel nucleic acid containing a specific segment having at least one region that modulates expression of the VRI (vanilloid receptor type 1) receptor, or a functional derivative, allele or fragment of this region, or a sequence that hybridises to it under standard conditions. The VRI modulator is derived from one or more of positions 221931-22334 of GenBank AL670399, 31673-36359 of AL663116, or 44731-43231 or 36616-33151 of AF188787 and is involved in transmission of pain, particularly in primary sensory neurons. The invention also describes a vector that contains the VRI modulator, host cells containing this vector (other than human germ or embryonal stem cells) and a method for modulator expression of the VRI receptor by introducing the condulator or the vector into a cell that contains the VRI gene. The products of the invention are used for detecting a transcription factor from its binding to a regulatory sequence (or a double-stranded collipson that modulates VRI receptor expression in cludes a sasociated with overexpression or underexpression of the transcription factor. The region that modulates VRI receptor expression includes a binding site for a transcription factor, e.g. MZEI, NEWappaB, NEAT or care used for preferred them, receptor expression includes a containing them,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PCR
 are used for prevention or treatment of pain, also for treating sensitivity disorders, e.g. analgesia, hypalgesia or homosomers, e.g. analgesia, hypalgesia or homosomers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             New nucleic acid that modulates expression of the vanilloid receptor-1, useful for control of pain or sensitivity disorders, comprises sequences from control regions of the receptor gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human VR1 exon 1d transcription factor binding fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADQ30417 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 54; 68pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          09-DEC-2002; 2002DE-01057421
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-DEC-2003; 2003WO-EP013522
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      hypalgesia; hyperalgesia; neuralgia; myalgia; human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      pain transmission; primary sensory neuron; transcription factor;
detection; MZFI; NFkappaB; NFAT; GATA1; sensitivity disorder; an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 09-SEP-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (CHEF )
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           VR1 receptor; vanilloid receptor type 1; modulator;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    primer which was
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2004-468868/44.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             859 AAAGTGCTGGGATTACAGGCGTGA 882
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GRUENENTHAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    24 BP; 7 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAAGTGCTGGGATTAGAGGCGTGA 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bieller A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2.3%;
95.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GMBH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2 C; 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Schaefer MKH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 22.4;
Pred. No. 9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    G; 5 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               9.7e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                also
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RESULT 359
ABX15537
ID ABX155
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                                                                                                                                                                                                                                                                                                                                                                                                                       watherosclerosis; autolimmune carditis; cardiomyopathy; ulcerative colitis; KW cardiac cell disfunction; aortic smooth muscle cell activation; trauma; KW cardiac cell apoptosis; gastrointestinal inflammation; cerebral trauma; KW cardiac cell apoptosis; gastrointestinal inflammation; cerebral trauma; KW Kawasaki's syndrome; cervical lymphadenopathy; diabetic nephropathy; KW Kawasaki's syndrome; cervical lymphadenopathy; diabetic nephropathy; KW glomerulonephritis; diabetic retinopathy; Grave's opthalmopathy; KW chronic lung disease; chronic sinusitis; chronic lymphocytic thyroditis; KW chronic plutic pain syndrome; alopecia areata; Grave's disease; KW thyroid disease; goiter; struma lymphomatosa; sleep disorder; neoplasia; KW chronic fatigue syndrome; obesity; infectious disease; Leishmaniasis; KW hoprosy; myocardial disfunction; breast cancer; organ transplant; KW Hodgkin's disease; hormonal regulation; fertility; septicaemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; ss; PCR; primer; interleukin-1; II-1; marker gz51gz6; nephropathy; inflammatory disease; Systemic Inflammatory Response; SIRS; genotyping; Alzheimer's disease; arthritis; acute joint inflammation; opthalmopathy; juvenile chronic arthritis; asthma; bronchial asthma; pulmonary disease; chronic obstructive airways disease; cardiovascular disease; thyroditis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 receptor. This sequence represents a fragment of human VR1 which is capable of binding to a transcription factor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABX15537;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABX15537 standard;
                                                                                                                                                                                                                                                                                 misc_difference
                                                                                                                                                                                                                                                                                                                                       modified_base
                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
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                                       WPI; 1999-080814/07.
                                                                                              (INTE-)
                                                                                                                          29-MAY-1997;
30-JUN-1999;
                                                                                                                                                                     27-APR-2001; 2001US-00845129
                                                                                                                                                                                               10-OCT-2002
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                                                                                              INTERLEUKIN GENETICS
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                                                                                                                           97GB-00011040
99US-00345217
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/mod_ba
                                                                                                                                                                                                                                                      /note= "Given in the specification as the number
                                                                                                                                                                                                                                                                                                 /note= "OTHER= FAM labelled"
                                                                                                                                                                                                                                                                     *tag=
                                                                  Camp
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95.8%;
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                                                                   D.
                                                                                               INC
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Pred. No. 9
                                                                   Giovine
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                                                                   FS
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New method of determining a patient's susceptibility to inflammatory disorders - by detecting the presence of an IL-1 (44112332) haplotype, $\vec{\imath}$

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CC determining or characterising the causative genetic variation. Diseases CC such as inflammatory disease e.g. Systemic Inflammatory Response (SIRS), CC Alzheimer's disease; arthritis e.g. acute joint inflammatory, Duvenile CC chronic arthritis; asthma e.g. bronchial asthma, chronic obstructive CC carditis; cardiomyopathy and cardiac cell disfunction e.g. autroimmune CC carditis; cardiomyopathy and cardiac cell disfunction e.g. acutic smooth CC muscle cell activation, cardiac cell apoptosis; gastrointestinal CC inflammations, inphropathies e.g. diabetic nephropathy, coronary CC infection; Kawasaki's syndrome e.g. cervical lymphadenopathy, coronary CC artery lesions; nephropathies e.g. diabetic nephropathy, Cronic glomerulonaphritis; opthalmopathies e.g. diabetic retinopathy, Grave's CC periodontal disease; pulmonary diseases e.g. chronic lung disease, CC chronic sinusitis; thyroditis e.g. chronic lymphacytic thyroditis; CC periodontal disease; pulmonary diseases e.g. chronic lung disease; CC urinary tract infections e.g. chronic prostatitis, chronic pelvic pain CC syndrome; immunological disorders e.g. alopecia areata, Graves disease; CC chronic fatigue syndrome; obesity; infectious disease e.g. Leprosy, CC chronic fatigue syndrome; obesity; infectious disease e.g. Leprosy, CC cishmaniasis; trauma e.g. cerebral trauma, myocardial disfunction; e.g. chronic palvasias e.g. chronic palvasias e.g. trauma e.g. cerebral trauma, myocardial disfunction; e.g. chronic activatis disease; hormonal regulation e.g. cerebral trauma, myocardial disfunction; e.g. chronic palvasias e.g. therefore, condition, e.g. a clinician can more effectively prescribe a therapy chart will address the molecular basis of the disease or condition. The present scripts for impers #1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 360
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     with an IL-1 inflammatory haplotype. The method involves detecting at least one allele of the haplotype, where the presence of the allele indicates that the subject is predisposed to the development or has the disease or condition. The invention allows the determination of an individual's likelihood for developing a particular disease or condition associated with interleukin 1 (IL-1) polymorphisms without necessarily
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to a method for determining whether a subject has or is predisposed to developing a disease or condition that is associated with an IL-1 inflammatory haplotype. The method involves detecting at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 5;
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                                                                                                                                                        Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                              SNP specific SNPE primer SEQ ID 1243.
                                                                                                                                                                                                                                                                                                                                           14-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                      AAH38447 standard; DNA; 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    marker gz51gz6 primer #1
                                                                                          Homo sapiens
                                                                                                                                        nflammation;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GGGATTACAGGCGTGAGCCACCACG 891
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                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                        forensic
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                                                                                                                                        investigation; paternity
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Pred. No. 1e
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CC SNP flanking sequence, the SNPE primer is used as a genotyping primer. CC The oligonucleotides are useful for genotyping a nucleic acid sample by performing a single-nucleotide primer extension reaction. The coligonucleotides are useful for determining the presence, absence or coligonucleotides are useful for determining the presence, absence or cc assess by association analysis the genotype of an individual or group of cindividuals, having a pathological phenotype trait suspected of being caused by one or more SNPs. Phenotypic traits include diseases e.g. cc agammaglobulinaemia, diabetes insipidus, Lesch-Nyhan syndrome, muscular cdystrophy, familial hypercholesterolaemia, polycystic kidney disease, cc osteogenesis imperfecta and acute intermittent porphyria. Phenotypic craits also include symptoms of or susceptibility to multifactorial confiammation, cancer, nervous system diseases and infection by pathogenic microorganism. The method is also useful in forensic investigations and conternity analysis. The present sequence represents a single nucleotide primer extension (SNPE) primer specific for a human SNP containing DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
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Best Local (
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                                                                                                                                                                                Claim 1; Page 56; 83pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Picoult-Newburg L,
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                                                                                                                                                                                                                                                                                                                                       Human MDZ7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADB04744 standard; DNA;
                                                                                                                         Homo sapiens
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                                                                                                                                                                                                                                                                                                                                   scanning oligonucleotide SEQ ID
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                                                                                                                                             5730
                                                                                                                                                                                                                      chromosome 15q26.1;
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Best Local S
Matches 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is encoded at chromosome 7g22.1, MDZ4 is encoded at chromosome 6p21.3-22.2, MDZ7 is encoded at chromosome 16p11.2 and MDZ12 is encoded at chromosome 15p11.2 and MDZ12 is encoded at chromosome 15p11.2 and MDZ12 is encoded at chromosome 15p26.1. The MDZ3, MDZ4, MDZ7, and MDZ12 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12 genetic locus. The probes are
Shannon M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ7 or MDZ12, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ7; MDZ1; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; developmental disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADB04742 standard; DNA; 25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 8; SEQ ID NO 5730; 103pp; English.
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                                                                                                                 02-AUG-2001; 2001US-00922181
                                                                                                                                                                            30-JUL-2002; 2002EP-00016874.
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                                                                                                                                                                                                                                                                                              EP1281758-A2
                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human MDZ7
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                                                          (AEOM-) AEOMICA INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25 BP; 5 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TTTGTATTTTAGTAGAGACGGGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TTTGTATTTTAGTAGAGATGGGG 794
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                scanning oligonucleotide SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GuY,
ပ်
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2001US-00922181
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               present sequence was used to illustrate the
Ķ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nguyen
Nguyen C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           eased or increased expression e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 C; 9 G; 10 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pred. No. 1e+00; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 22.4;
Pred. No. 1e
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4; DB 1;
1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    5728
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                  cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0
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profeins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is encoded at chromosome 7922.1, MDZ4 is encoded at chromosome 6p21.3-22.2, MDZ7 is encoded at chromosome 15p11.2 and MDZ12 is encoded at chromosome 15p11.2 and MDZ12 is encoded at chromosome 15q26.1. The MDZ3, MDZ4, MDZ7, and MDZ12 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for dispnosing or monitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12 genetic locus. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ4, MDZ7 or MDZ12, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention relates to novel human zinc finger-containing profeins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 8; SEQ ID NO 5728; 103pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2003-423107/40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MDZ3,
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Query Match Best Local 9 Matches Sequence 25 770 TTTTGTATTTTTAGTAGAGATGGG 793 23; Similarity BP; 6 A; 1 C; 7 G; 11 T; 0 U; 2.3%; llarity 95.8%; Conservative 0 Pred. No. le+0 0; Mismatches Score 22.4; Pred. No. 16 1; DB 1; Length 25; 1e+03; 0 Gaps

0

0 Other;

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RESULT 363
ADO12082/c
ADO12082 standard;
                                                          15-JUL-2004
                                                              ADO12082;
                                                          (first entry)
                                                                  DNA;
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N

TTTTGTATTTTAGTAGAGACGGG

Single multiplex PCR

primer #1454.

ss; primer; simultaneous amplification; single multiplex polymerase chain reaction; multifactorial disease; genetic alteration; pharmacogenetic reaction; genotyping; polymorph gene expression profiling. Synthetic genotyping; polymorphism;

WO2004033649-A2

22-APR-2004

07-OCT-2003; 2003WO-US031874

07-OCT-2002; 2002US-0417009P

AIND NEW JERSEY MEDICINE & DENTISTRY

Li H, Ŀ

WPI; 2004-340914/31.

Designing primers for simultaneous amplification of target DNA fragments in a single multiplex polymerase chain reaction, for high throughput multiplex DNA sequence amplification, comprises aligning two primers.

Disclosure; Page 39; 120pp; English

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CC amplification relates to a method of designing primers for simultaneous cc amplification of target DNA fragments in a single multiplex polymerase cc chain reaction by aligning a first primer and a second primer. The method cc comprises (a) aligning a first primer and a second primer, and (b) cc selecting the first primer where the first primer at its 3' end does not contain four or more bases that are perfectly matching to the 3' end cc sequence of the first primer at at its 3' end does not contain seven or more bases that are perfectly matching to the 3' end cc second primer, the first primer at its 3' end does not contain six or contain six or contain six or contain six or the second primer at the first primer at its 3' end does not contain six or contain six or the second primer, the first primer at its 3' end consess that are perfectly matching to a sequence anywhere of the first primer or the second primer, and the first primer at its 3' end con sismatch to a sequence anywhere of the first primer or the second contain exaction. It is also useful for designing primers for simultaneous amplification. It is also useful in the identification of multiple genes chain reaction, the studies in pharmacogenetic reactions, the genotyping conservance contains in a large population, the gene expression contains in a large population, the gene expression this sequence corresponds to an example of a primer of the invention.
RESULT 364
ADO12035
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ij
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local S
Matches 24
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                                                                  Designing primers for simultaneous amplification of target DNA fragments in a single multiplex polymerase chain reaction, for high throughput multiplex DNA sequence amplification, comprises aligning two primers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ss; primer; simultaneous amplification; single multiplex polymerase chain react
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADO12035 standard; DNA; 27
                                                                                                                                                                                                                                                                                        07-OCT-2003; 2003WO-US031874.
                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Single multiplex PCR primer #1407
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15-JUL-2004
                                                                                                                                           WPI; 2004-340914/31.
                                                                                                                                                                                                                    (UYNE-) UNIV NEW JERSEY MEDICINE & DENTISTRY
                                                                                                                                                                                                                                                        07-OCT-2002; 2002US-0417009P
                                                                                                                                                                                                                                                                                                                               22-APR-2004
                                                                                                                                                                                                                                                                                                                                                                 WO2004033649-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                        expression
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               24;
                                                                                                                                                                                 댰
                                                                                                                                                                                                                                                                                                                                                                                                                                      multiplex polymerase chain reaction; multifactorial disease; alteration; pharmacogenetic reaction; genotyping; polymorph pression profiling.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP; 5 A; 6 C; 11 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ВP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 22.2;
Pred. No. 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             genotyping; polymorphism;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 27;
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Disclosure; Page 39; 120pp; English

relates

ç g)

method of

designing

primers for simultaneous

New tumourigenesis T-genes and proteins - useful for, antibodies for clinically diagnosing cells having non-

This is the nucleotide sequence of amplification in the method of the

the PCR primer P2 used for invention, which involves isolation

e f

amplification

Example

Page

6

71pp;

English.

capacity

such

diagnosing cells ha ch as lipoblastomas.

for, e.g. preparing non-physiological

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AAV29285
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cc amplification of target DNA fragments in a single multiplex polymerase cc chain reaction by aligning a first primer and a second primer. The method cc comprises: (a) aligning a first primer and a second primer; and (b) cc selecting the first primer where the first primer at its 3' end does not contain four or more bases that are perfectly matching to the 3' end cc sequence of the first primer or a second primer, the first primer at its 3' end does not contain seven or more bases that are perfectly matching cc except one mismatch to the 3' end sequence of the first primer or the second primer, the first primer at its 3' end does not contain six or cmore bases that are perfectly matching to a sequence anywhere of the first primer or the second primer, and the first primer at its 3' end does not contain six or contain seven or more bases that are perfectly matching except one mismatch to a sequence anywhere of the first primer or the second cc primer. The method is useful for designing primers for simultaneous camplification of target DNA fragments in a single multiplex polymerase chain reaction. It is also useful in the identification of multiple genes related to multifactorial diseases, the genome-scale detection of genetic calterations, the studies in pharmacogenetic reactions, the genotyping confiling in various samples and high throughput genotyping technologies. This sequence corresponds to an example of a primer of the invention.
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RESULT 365
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 27 BP; 5 A; 11 C; 6
                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; tumourigenesis gene; T-gene; PLAG2; PLAG1; CTNNB1; antibody; benign tumour; malignant tumour; leukaemia; lymphoma; cancer; inhibition;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAV29285 standard; cDNA; 22
                                                                                                                                                                                                                                                                                                                                             EP825198-A1
                                                                                                                                                                                                                                                                                                                                                                                           Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21-AUG-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAV29285;
                                                                                                                                         WPI; 1998-132252/13.
                                                                                                                                                                     Van De Ven WJM,
                                                                                                                                                                                                                                                 22-AUG-1996;
                                                                                                                                                                                                                                                                                17-JAN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                          PCR; amplification; primer;
                                                                                                                                                                                                  (KULE-) KU LEUVEN RES &
                                                                                                                                                                                                                                                                                                               25-FEB-1998
                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          661 GGCGCAATCTTGGCTCACTGCAACCTC 687
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            24;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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                                                                                                                                                                                                                                                                               97EP-00200130
                                                                                                                                                                                                                                                   96EP-00202339
                                                                                                                                                                      Stenman KGD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        of,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PCR primer
                                                                                                                                                                                                    DEV.
HOLDINGBOLAGET AB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 22.2;
Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          G; 5 T; 0 U;
                                                                                                                                                                        Kas KP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ρ2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   27
                                                                                                                                                                        Voz ML
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                27;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
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និក្សាស្ត្រាស់និងស្ត្រាស់ស្ត្រ ស្ត្រាស់ស្ត្រ ស្ត្រ ស្ត្រាស់ស្ត្រ ស្ត្រ ស្ត្រ ស្ត្រ ស្ត្រ ស្ត្រ ស្ត្រាស់ស្ត្រ ស្ត្រ ស្ត្រ

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Matches
cows. The protein is expressed and secreted into the milk as a fusion protein that has reduced biological activity. The animal is then milked and the fusion protein is then cleaved (chemically or enzymatically) to release the desired active protein which may then be isolated and utilized. The use of to recombinantly produce polypeptides in milk minimizes health problems in the animal and prevents side effects associated with ectopic expression or leakage of the protein into surrounding tissues and the circulation. These side effects are a particular risk when producing potent polypeptides such as growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 the tumourigenesis genes (T-gene), in the form of PLAG1, PLAG2, and CTNNB1 genes. Their proteins can be used as a starting point for preparing antibodies for clinically/medically diagnosing cells having a non-physiological proliferative capacity as compared to wild type cells, where the former cells are selected from both benign and malignant tumours, as well as leukaemia and lymphomas. Derivatives of the T-gene are also used in the diagnosis and preparation of therapeutical compositions for the treatment of cancers, such as nucleic acid derivatives, and antibodies. The T-gene may be uses as a starting point for designing suitable expression-modulating compounds or techniques for the treatment of non-physiological proliferation phenomena in humans or animals. Expression inhibitors of the T-gene can be used in the treatment
                                                                                                                                                                                                                                                                                                        The invention relates to a new process for producing biologically active polypeptides (e.g. erythropoletin (EPO)) in mammalian milk as fusion proteins that are less active (or non-active) than the free polypeptides. The process may be used for the recombinant expression of proteins (especially EPO) in the milk of transgenic animals such as sheep and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 1; Col 7; 10pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Producing biologically active polypeptides in mammalian milk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17-AUG-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Erythropoietin; EPO; mammalian milk; transgenic animal; lactation; ectopic expression; growth factor; cytokine; enzyme; transgene; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AHRECASEPO transgene
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (PHAR-) PHARMING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            385
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hyttinen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative (
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RESULT 367
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                                                                                                                   The invention relates to human chromosomal region 9q11 (AAC87588). Abnormalities in this region of the short arm of chromosome 9 is thought to be associated with miscarriage and cancer, as an ovarian cancer patient with a history of miscarriage was found to have a chromosomal inversion inv(9) (p11;q13). The 9p11 region contains the gene encoding keratinocyte growth factor (KGF), and the invention also specifically claims the KGF PCR primers AAC87589 and AAC87590 for use in detecting all or part of the KGF gene. The nucleic acid sequences can be used to detect abnormalities in chromosomal region 9p11 and thus give an indication of an individual's risk of developing a 9p11-associated condition. Sequences AAC87596-C87597 represent human Alu sequence PCR primers used in an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   factors, cytokines and enzymes. This means the transgenic animal remains healthy, viable and able to lactate for longer. Sequences AAZ07499-500 represent AHRECASEPO transgene specific primers used for screening GO mice. AHRECASEPO is a gene construct designed to secrete biologically active free human EPO in transgenic mouse milk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; keratinocyte growth factor; KGF; chromosome cancer; miscarriage; spontaneous abortion; genetic diagnosis; Alu sequence; PCR primer; ss.
                                                                     Sequence
                                                                                                       exemplification fo the invention
                                                                                                                                                                                                                                                                                                                                                                                                     The base
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                 Local
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l Similarity
22; Conser
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larity 100.0%;
Conservative
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                                                                       G; 5 T;
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. 9.6e+02;
9.6e+02;
                                   DB 1;
                                                                       0 Other;
                                  Length 22;
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susceptibility;
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RESULT 368
AAF88160
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Matches
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                                                                                                                                                                                                                                                                                                                                This invention describes a novel nucleic acid (NI) encoding a polypeptide which comprises a KRAB-domain and/or at least one zinc finger motif. The products of the invention have cytostatic and antithyroid activity and can be used in gene therapy. Nucleic acids, polypeptides, and antibodies of the invention may be used in the diagnosis and/or the therapy of the malfunction of the thyroid and/or hyperlasias of the thyroid and/or thyroid tumors. They may also be used in the production of medicaments. (NI) can also be used to diagnose thyroid tumors which are located on chromosome 19 at band 19q13. This sequence represents a PCR primer used in the isolation of the thyroid malfunction-associated protein, RITA which is described in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        KRAB domain; hyperplasia; thyroid; tumor; zinc finger motif; primer; cytostatic; antithyroid; gene therapy; chromosome 19; 19q13; ss.
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                                                                                                                                                                                                                                                                                                            Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New nucleic acid useful for the diagnosis and treatment of thyroid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bullerdiek J,
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                                Cytostatic; Antibacterial; Immunosuppressive; Antiinflammatory; neural thread protein; NTP; tumour; ds.
                                                                                                                                             ACC84464 standard; DNA; 22 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (UYBR-) UNIV BREMEN.
                                                                      NTP peptide
                                                                                               28-AUG-2003
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                                                                     encoding sequence #11.
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                                                                                               (first entry)
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                                                                                                                                                                                                                                                                                                              6 G;
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RESULT 370
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        antibacterial, immunosuppressive and antiinflammatory. It is useful for treating a condition in a patient requiring removal or destruction of cells, for treating a condition such as benign or malignant tumor, inflammatory disease, autoimmune disease and infectious disease. The peptide useful for treatment is derived from the amino acid sequence for a pancreatic thread protein. The peptide is conjugated, linked or bound to a molecule chosen from antibody or its fragment, antibody-like binding molecule, where the molecule has a higher affinity for binding to a tumor or other target than binding to other cells. Treatment using NTP peptides can remove benign tumors with less risk and fewer of the undesirable side effects of surgery. The present sequence is an NTP encoding sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel neural thread protein peptide, referred as cell death peptide, useful for treating prostatic hyperplasia, psoriasis, eczema, dermatosis, atherosclerosis, cosmetic modification to skin, throat, mouth, muscle.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19-JUL-2001; 2001US-0306150P.
19-JUL-2001; 2001US-0306161P.
16-NOV-2001; 2001US-0331477P.
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                                                                                                                                                    Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to a neural thread protein (NTP) referred to as cell death peptide. Thought to be cytostatic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19-JUL-2002; 2002WO-CA001105
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                                                                                                                                                                                Mouse; survival SMA; diagnosis;
                                                                                                                                                                                                                          Mouse SMN 5'
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                                                                                                                                                                                                                                                                                  AAH44054;
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              (SINI-) ACAD SINICA
                                                                    25-MAY-2000; 2000US-00578656
                                                                                                12-JUN-2001.
                                                                                                                           US6245963-B1
                                         28-MAY-1999;
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                                                                                                                                                                                                                          untranslated region PCR primer SEQ ID
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                                         99US-0136520P
                                                                                                                                                                                   detection;
                                                                                                                                                                                               motor neuron; SMN; knockout; spinal muscular atrophy;
                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%; F1
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9.6e+02;
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AAH38407
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Best Local Similarity
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New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nucleic
                                                                                                                                                                                                                                                                                                                                      Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genoryping; agammaglobulinaemia; diabetes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                          SNP specific SNPE primer SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAH38407 standard; DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention describes a transgenic mouse whose genome comprises a homozygous disruption of an Smn (survival motor neuron) gene which does not produce functional Smn protein. The mouse genome additionally comprises a DNA sequence encoding human SMN protein, where expression of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           muscular atrophy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New transgenic mouse having a genome comprising a homozygous disruption of an Smn gene, useful e.g. as a model for human spinal muscular atrophy, or for testing the efficacy of present or future treatments for spinal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             닭
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                                                                                                                                                      15-OCT-1999;
                                                                                                                                                                                    13-OCT-2000; 2000WO-US028436.
                                                                                                                                                                                                                       26-APR-2001
                                                                                                                                                                                                                                                                                          Homo
                                                                                                                                                                                                                                                                                                                         inflammation;
                                                    2001-290930/30
                                                                                                                                                                                                                                                                                          sapiens
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                                                                                                                     ORCHID
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ilarity 100.0%;
Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                           forensic
                                                                                                                     BIOSCIENCES INC
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                                                                                    Pohl
                                                                                                                                                                                                                                                                                                                           investigation;
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Pred. No.
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                                                                                                                                                                                                                                                                                                                        paternity
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1e+03;
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CC includes kits for determining the presence of a SNP, using the coligonucleotides of the invention. The PCR primers are used to amplify a CC SNP flanking sequence, the SNPB primer is used as a genotyping primer. CC The oligonucleotides are useful for genotyping a nucleic acid sample by CC performing a single-nucleotide primer extension reaction. The CC oligonucleotides are useful for determining the presence, absence or CC oligonucleotides are useful for determining the presence, absence or CC identity of a SNP and for genotyping nucleic acid samples, for e.g. to CC assess by association analysis the genotype of an individual or group of CC individuals, having a pathological phenotypic trait suspected of being CC caused by one or more SNPs. Phenotypic traits include diseases e.g. CC agammaglobulinaemia, diabete sinsipidus, Lesch-Nyhan syndrome, muscular CC diseases imperfects and caute intermittent porphyria. Phenotypic CC osteogenesis imperfects and caute intermittent porphyria. Phenotypic CC disease of which a component is or may be genetic such as autoimmune CC diseases, including, rheumatoid arthritis, multiple sclerosis, pathogenic confirmmation, cancer, nervous system diseases and infection by pathogenic confirmment of also inseril in forencie investications and
                                                                             microorganism. The method is also useful in forensic investigations and paternity analysis. The present sequence represents a single nucleotide primer extension (SNPE) primer specific for a human SNP containing DNA
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  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide primer extension (SNPE) primers, and the sequences of regions flanking
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            acid sample.
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25 BP; 6
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     7
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  6 T;
  0 U;
     1 Other;
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Query Match Best Local Matches 724 TCCTGAGTAGCTGGGACTACAGGC 747 l Similarity 91.7 22; Conservative TCCTRAGTAGCTGGGATTACAGGC 24 2.2%; 91.7%; Score 22; Pred. No. Mismatches 1e+03; DB 1; Length 25; Indels <u>,</u> Gaps 0

RESULT 372 ADB04745 20-NOV-2003 Human MDZ7 scanning oligonucleotide SEQ ID ADB04745 standard; DNA; (first entry) 25 ВÞ 5731.

Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ7; MDZ12; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 15p11.2; chromosome 15q26.1; cancer; developmental disorder; ss.

Homo sapiens

EP1281758-A2

05-FEB-2003

30-JUL-2002; 2002EP-00016874

02-AUG-2001; 2001US-00922181

Z, ည Y, Nguyen

2003-423107/40

New zinc finger-containing manufacturing a medicament proteins and for treating nucleic acids, useful or preventing a disord

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KW Zinc 1
KW Chroma
KW EP1281
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ7; MDZ12; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human MDZ7 scanning oligonucleotide SEQ ID 5565
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADB04579 standard; DNA; 25 BP
The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is encoded at chromosome 7q22.1, MDZ4 is encoded at chromosome 6p21.3-22.2,
                                                                                                                                                                                                               New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ7 or MDZ12, e.g. cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Shannon M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             02-AUG-2001; 2001US-00922181
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                                                                                                                                                Example 8;
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                                                                                                                                                SEQ ID NO 5565; 103pp; English.
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Pred. No. 1
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Best Local 9
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The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is encoded at chromosome 7g22.1, MDZ4 is encoded at chromosome 6p21.3-22.2, MDZ7 is encoded at chromosome 1fp11.2 and MDZ12 is encoded at chromosome 1fp11.2 and MDZ12 is encoded at chromosome 1fp11.2 and MDZ12 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ7; MDZ12; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1;
                                                                                                                                                                                                                                                                            New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ4, MDZ7 or MDZ12, e.g. cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADB04741 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                        Shannon M,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                developmental disorder; ss.
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                                                                                                                                                                                                                                         SEQ ID NO 5727; 103pp; English.
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Pred. No. 1.
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alterations in MDZ3, MDZ4, MDZ7,

or MDZ12 genetic locus. The probes are

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ADJ33167/c
ID ADJ331
AC ADJ31
AC ADJ3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Antiinflammatory; nephrotropic; hepatotropic; neuroprotective; nootropic; gynaecological; cytostatic; antiallergic; immunosuppressive; antithyroid; antiparkinsonian; antiarthritic; monocarboxylic acid; transport protein; inhibitor; potentiator; organic ion; TCH131; TCH182; TCH120; respiratory disease; asthma; kidney disease; kidney failure; nervous system disease; Alzheimer's disease; muscle disease; muscle wasting; allergic disease; meningitis; autoimmune disease; multiple sclerosis, allergic disease; hayfever; spleen disease; immune deficiency disease; leukopenia; liver disease; hepatitis; digestive disease; Crohn's disease; genital disease; ovarian hypofunction; cancer; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention.
The invention relates to proteins TCH131 (human, mouse and rat), TCH182 (human) and TCH120 (human) and their salts and partial peptides, and similar proteins with equivalent activity. Also disclosed are polynucleotides (including DNA) encoding the proteins. Proteins of the invention are useful in the prevention, treatment and diagnosis of respiratory diseases (including asthma and bronchitis), kidney diseases (including kidney failure and nephritis), nervous system diseases (including Alzheimer's, Parkinson's and schizophrenia), metabolic acidosis, muscle diseases (including muscle wasting), allergic diseases (including pneumonia, meningitis and myocarditis), autoimmune diseases
                                                                                                                                                                                                                                                                                                                                                                                                              Monocarboxylic acid and organic ion transport proteins and compounds modifying their activity or expression for treatment, prevention and diagnosis of respiratory, inflammatory, autoimmune, allergic and kiddiseases and cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     07-NOV-2001; 2001JP-00342139
16-NOV-2001; 2001JP-00351086
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The present invention describes a method for detecting the presence polymorphisms associated with inflammatory bowel diseases such as ulcerative colitis and Crohn's disease. The methods can be used to d the presence of genetic polymorphisms associated with inflammatory be disease and correlating their occurrence with disease states. They used in this way for phenotypic correlations, forensics, paternity testing, medicine and genetic analysis. The present sequence is a polymorphic site described in the exemplification of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; inflammatory bowel disease; Crohn's disease; ulcerative colitis; single nucleotide polymorphism; SNP; chromosome 19p13; paternity test; chromosome 5q31-33; forensic test; gene therapy; ds.
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                                                                                                                                           The present invention describes a method for detecting the presence of polymorphisms associated with inflammatory bowel diseases such as ulcerative colitis and Crohn's disease. The methods can be used to detect the presence of genetic polymorphisms associated with inflammatory bowel disease and correlating their occurrence with disease states. They may be
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                                                                                                                                                                                                                  Page 46; 463pp; English.
                                                                                                                                                                                                                                                                                                 Hudson TJ,
                     GTGCTGGGATTACAGGCGTGAGCCAC 887
 GTGCTGGGATTGCANGTGTGAGCCAC 1
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                                                      2.2%;
                                                                                         10 C;
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Pred.
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Pred. No. 1.1e
0; Mismatches
                                                                                         G; 4 T; 0 U;
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                                                                                                                                                                                                                                                                                                  Rioux J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21.8;
No. 1
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                                                        .1e+03
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                                                                   DB 1;
                                                                                          1 Other;
                                                                                                                                                                                                                                                                                                   Siminovitch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length
                                                                  Length 26;
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                                                                                                                                     paternity
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671 TGGCTCACTGCAACCTCTGCCTCCCGG 697

Query Match Best Local S Matches 23

Similarity

2.2%; llarity 85.2%; Conservative

Score 21.8; D. Pred. No. 1.1e. 0; Mismatches

.1e+03

4,

0

Gaps

0

DB

Length Indels

27;

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RABSULT 378
AAH38507/c
ID AH3855
XX AAH3855
XX AAH3855
XX AAH3855
XX Single
XX WO2001
XX Single

                                                                                                                                                                                                                    Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide primer extension (SNPE) primers, and the sequences of regions flanking controlled includes kits for determining the present or absence of a SNP, using the coligonucleotides of the invention. The PCR primers are used to amplify a CC SNP flanking sequence, the SNPE primer is used as a genotyping primer. CC The oligonucleotides are useful for genotyping a nucleic acid sample by performing a single-nucleotide primer extension reaction. The CC oligonucleotides are useful for determining the presence, absence or CC identity of a SNP and for genotyping nucleic acid sample by collective and for genotyping nucleic acid samples. For e.g. to assess by association analysis the genotype of an individual or group of claused by one or more SNPs. Phenotypic trait suspected of being cases by an additionally disbets insipidus, Lesch-Nyhan syndrome, muscular dystrophy, familial hypercholesterolaemia, polycystic kidney disease, considerable and acute intermittent porphyria. Phenotypic traits also include symptoms of or susceptibility to multifactorial diseases, including, rheumatoid arthritis, multiple sclerosis, confidence in revusa system disease and infection by pathogenic confidence in concern revous system disease and infection by pathogenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia, polycystic kidney disease; osteogenesis imperfecta; autoimmune disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAH38507;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     14-AUG-2001
                                                                                            inflammation, cancer, nervous system diseases and infection by pathogenic microorganism. The method is also useful in forensic investigations and paternity analysis. The present sequence represents a single nucleotide primer extension (SNPE) primer specific for a human SNP containing DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200129262-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Picoult-Newburg L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13-OCT-2000; 2000WO-US028436
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        inflammation;
   Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2001-290930/30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        stic kidney disease; osteogenesis imperfecta; autoimmune disease; intermittent porphyria; rheumatoid arthritis; multiple sclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1; Page 56; 83pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORCHID BIOSCIENCES INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            standard; DNA;
       27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ping; agammaglobulinaemia; diabetes insipidus; cancer; syndrome; muscular dystrophy; familial hypercholesterolaemia;
       B₽;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               99US-0160096P.
       8
       A;
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       6 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            investigation; paternity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   27
          8
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              ς,
              2 Other;
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TGGCTCACTGNAACCTCTGNCTTCTGG 1

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RESULT 379
AAH37975/c
                                                                              Query Match
Best Local S
Matches 23
                                                                                                                                                                                                                                                                                                                  SNP flanking sequence, the SNPE primer is used as a genotyping primer. The oligonucleotides are useful for genotyping a nucleic acid sample by performing a single-nucleotide primer extension reaction. The oligonucleotides are useful for determining the presence, absence or identity of a SNP and for genotyping nucleic acid samples, for e.g. to assess by association analysis the genotype of an individual or group of individuals, having a pathological phenotypic trait suspected of being caused by one or more SNPs. Phenotypic traits include diseases e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 7975/c
AAH37975
                                                                                                                                                                                                        agammaglobulinaemia, diabetes insipidus, Lesch-Nyhan syndrome, muscul dystrophy, familial hypercholesterolaemia, polycystic kidney disease, osteogenesis imperfecta and acute intermittent porphyria. Phenotypic traits also include symptoms of or susceptibility to multifactorial disease of which a component is or may be genetic such as autoimmune diseases, including, rheumatoid arthritis, multiple sclerosis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo
                                                                                                                                 inflammation, cancer, nervous system diseases and infection by pathogenic microorganism. The method is also useful in forensic investigations and paternity analysis. The present sequence represents a single nucleotide primer extension (SNPE) primer specific for a human SNP containing DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            primer extension (SNPB) primers, and the sequences of regions flanking sites of single nucleotide polymorphisms SNPs. The present invention includes kits for determining the presence or absence of a SNP, using the coligonucleotides of the invention. The PCR primers are used to amplify a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; Page 53; 83pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Picoult-Newburg L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     26-APR-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SNPE; genotyping; agammaglobulinaemia; dlabetes insipidus; cancer; lesch Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis; inflammation; forensic investigation; paternity analysis; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Single nucleotide polymorphism; SNP; single nucleotide primer extension;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SNP specific SNPE primer SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15-OCT-1999;
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                     Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sample.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 standard; DNA; 27
     Conservative
                                                                            BP; 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
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                   2.2%;
                                                                            6 C; 8 G; 5 T; 0 U;
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   <u>.</u>
                   Score 21.8;
Pred. No. 1.
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     Mismatches
                                                                            2 Other;
                     .le+03
                                       DB 1; Length
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Gaps
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RESULT 381
AAX83037/c
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AAH91552
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                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; inflammatory bowel disease; Crohn's disease; ulcerative colitis single nucleotide polymorphism; SNP; chromosome 19p13; paternity test; chromosome 5q31-33; forensic test; gene therapy; ds.
                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                         the presence of genetic polymorphisms associated with inflammatory disease and correlating their occurrence with disease states. They used in this way for phenotypic correlations, forensics, paternity testing, medicine and genetic analysis. The present sequence is a polymorphic site described in the exemplification of the invention
  AAX83037
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention describes a method for detecting the presence of polymorphisms associated with inflammatory bowel diseases such as ulcerative colitis and Crohn's disease. The methods can be used to detect the methods can be used to detect
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Testing for the presence of polymorphisms associated with inflammatory bowel disease, using a hybridization assay.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2001-367874/38.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Daly M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (ELLI-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                10-DEC-1999; 99US-0170257P.
10-APR-2000; 2000US-0196046P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human inflammatory bowel disease associated polymorphic site #627.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           09-OCT-2001
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                                                                                                                                                                                                                                                                   Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    485
                                                                                                                                                                                382 GCCTCCCAAAGTGCTGGGATTACAGG 407
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                                                                                                                                                                                                                                          23;
                                                                                                                                                                                                                                                                   Similarity
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                                                                                                                                                                                                                                                                                                                                                27 BP; 7
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standard; DNA;
                                                                                                                                    GCCTTCCAAAGTGCNAGGATTACAGG
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                                                                                                                                                                                                                                            Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        note= "SNP, optionally T or C at this position"
                                                                                                                                                                                                                                                                                                                                                Α,
                                                                                                                                                                                                                                                             2.2%;
                                                                                                                                                                                                                                                                                                                                                6 C; 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Lander ES,
  23
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  ВP
                                                                                                                                                                                                                                          <u>,</u>
                                                                                                                                                                                                                                                                                                                                                G; 6 T; 0 U;
                                                                                                                                                                                                                                                                   Score 21.8;
Pred. No. 1
                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Rioux J,
                                                                                                                                    26
                                                                                                                                                                                                                                                                                                                                                1 Other;
                                                                                                                                                                                                                                                                     .1e+03;
                                                                                                                                                                                                                                                                                           DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Siminovitch
                                                                                                                                                                                                                                                                                         Length 27;
                                                                                                                                                                                                                                            Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  colitis;
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Best Local S
Matches 22
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30-JAN-1996;
12-APR-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Primer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Primers AAX83008-X83064 were used to RT-PCR amplify exons from the 5' and 3' ends of the human WRN gene (AAX83003) which encodes a protein related to Werner's syndrome. The products can be used for the detection and treatment of Werner's syndrome (WS), an autosomal recessive disorder with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  detection
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29-DEC-1995;
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                                                                                                                                                                                                        Human; zinc finger protein 15; cytostatic; virucidal; immunomodulatory; antiinflammatory; haemostatic; gene therapy; malignant neoplasm; haemopathy; HIV infection; immunological disease inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
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                                                                20-SEP-2001.
                                                                                                                                                                                     haemopathy;
PCR primer;
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                     26-FEB-2001; 2001WO-CN000165
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ive disorder; phenotype; primer; RT-PCR; amplification; ss
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96US-00594242.
96US-00632175.
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95US-00580539.
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                                                                                                                                                                                                                                                                                                                                                                                                                   DNA;
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Pred. No. 1.
                                                                                                                                                                                                                                                                                          PCR primer #1.
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and related diseases
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The present invention relates to the invention of the polynucleotide sequence encoding it described is the process for preparing the protein by DNA and the application of the polypeptide and polynucleotide

invention relates to the isolation of human zinc and the nolvnucleotide sequence encoding it. F

New human zinc finger protein 10.01 polypeptide for treating malignant tumors, hemopathy, human immunodeficiency virus infection, immunological

diseases and various inflammations.

Example

2; Page 17 (disclosure); 33pp; Chinese

WPI; 2002-692406/75.

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Xie BODE

(BODE-)

GENE

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COLTD

SHANGHAI

06-NOV-2000; 2000CN-00127241

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RESULT 383
ABS58183/c
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Matches 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human zinc finger protein 15 and encoded polynucleotide, appdiagnosis and treatment of malignant neoplasm, hemopathy, humannodeficiency virus infection, immunological diseases and
                                                                                                                                                                                                                                                                         Human; zinc finger protein 10.01; malignant tumour; haemopathy; human immunodeficiency virus infection; HIV infection; inflammation; immunological disease; RT-PCR; primer; reverse transcriptase-PCR; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention relates to human zinc finger protein 15 (s AAI65097 and AAG78828). The zinc finger protein and its coding s are useful in the diagnosis and treatment of malignant nepplasm, haemopathy, HIV infection, immunological diseases and various inflammations. The present sequence is a PCR primer, which was u example from the present invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 24 BP; 5 A; 5 C; 9 G; 5 T; 0 U;
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                                                                                                                                                                                                                                                                                                                   RT-PCR
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                                                                                                                                                                                                                                                      Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                           866 TGGGATTACAGGCGTGAGCCACC 888
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                                                                                                                                                                                                                                                                                                                  primer
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Matches 22
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Best Local &
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                                                                                                                                described is the process for preparing the protein by DNA recombination and the application of the polypeptide and polynucleotide in treating various diseases such as malignant tumours, hademopathy, human immunodeficiency virus (HIV) infection, immunological diseases, and various inflammations. The present sequence represents a reverse transcriptase (RT)-PCR primer used to isolate cDNA encoding human zinc
                                                                                                                                                                                                                                                                                                  New
                                                                                                                                                                                                                                                                                                                                                  Mao
                                                                                              Sequence
                                                                                                                                                                                                           The present invention relates to the isolation of human zinc finger protein 10.01, and the polynucleotide sequence encoding it. Also
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                                                                                                                      finger protein 10.01
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                                                           Similarity
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95.7%;
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                                                                        DВ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Other;
                                                                      <u>س</u>ر
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length
                                                                      Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           protein 10.01.
                                                                       24;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       human zinc
                                              0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               and
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RESULT 385
ABA04737
ID ABA047
XX ABA047
XX ABA047
XX ABA047
XX Human
XX Human;
KW Human;
KW Haemog
KW PCR pr
XX PCR PCR
XX PCR PCR
XX PCR
PN WO2001
XX PCR
PN W
                                                                                                                                                                                                                                                                                   RESULT 386
AAL45771/c
ID AAL457
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to human alkylation-DNA-protein cysteine methyltransferase (see AAM47739). The protein and its coding sequence as useful in the diagnosis and treatment of malignant tumours, haemopathy, HIV infection, immunological diseases and various inflammations. The present sequence is a PCR primer, which was used in an example from the
                                                                                                                                                                                                                              AAL45771;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human alkylation-DNA-protein cysteine methyltransferase and encoding polynucleotide, used in diagnosis and treatment of malignant tumors, hemopathy, human immunodeficiency virus infection, immunological diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; alkylation DNA protein cysteine methyltransferase 11; cytostatic; haemostatic; virucide; immunomodulatory; antiinflammatory; gene therapy;
                             Human; acid phosphatase family protein 11; cancer; haemopathy;
cytostatic; haemostatic; virucide; immunomodulatory; antiinflammatory;
                                                                                                                                                                        28-JUN-2002
                                                                                                                                                                                                                                                                                   AAL45771 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   and inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-055701/07.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28-MAR-2000; 2000CN-00115226
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   26-MAR-2001; 2001WO-CN000464.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO200188146-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human alkylation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABA04737;
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                                                                                                                   Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           22-NOV-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         767
                                                                                                                acid phosphatase family protein 11
  disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                 2 TTTTTTGTATTTTAGTAGTGA 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TTTTTTTGTATTTTTAGTAGAGA 789
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Page 19; 40pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP; 4 A; 0 C; 4 G; 16 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA
                                                                                                                                                                                                                                                                                         DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2.2%;
95.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 21.4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cysteine methyltransferase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pred. No. 1.1e+03;
  phlogosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                   CDNA
gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1;
                                                                                                                   PCR
therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length
                                                                                                                   primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PCR
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  primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sequence are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            primer #2
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Homo sapiens

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                                                                                                                                                                                                                                                                                                                                                              RESULT 387
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Мао У,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200220579-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention provides the protein and coding sequences of human acid phosphatase family protein 11. The sequences can be used in the treatment of cancer, haemopathy. HIV infection, immune diseases and phlogosis. The present sequence is a PCR primer for the coding sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21-JUN-2000; 2000CN-00116667.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 24 BP; 5 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19-JUN-2001; 2001WO-CN001011.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo acid phosphatase family protein 11 and encoding polynucleotide, in diagnosis and treatment of malignant tumors, hemopathy, human immunodeficiency virus infection, immunological diseases and
                  Reducing stutter in the amplification of a microsatellite for analysis by contacting the sample comprising a microsatellite enzyme with nucleic acid polymerase activity and incubating th with the enzyme.
                                                                                                                                     07-MAY-2001; 2001US-00850514.
                                                                                                                                                         06-MAY-2002, 2002WO-US014189
                                                                                                                                                                               14-NOV-2002
                                                                                                                                                                                                     WO200290562-A2
                                                                                                                                                                                                                                                         Stutter reduction;
                                                                                                                                                                                                                                                                             PCR primer #1 used to illustrate the method of the invention
                                                                                                                                                                                                                                                                                                  24-MAR-2003
                                                                                                                                                                                                                                                                                                                                           AAD50373
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (BIOW-)
                                                                                                                 (BIOW ) APPLIED BIOSYSTEMS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2002-329869/36.
                                                                         2003-111983/10.
                                                                                                                                                                                                                                                                                                                                                                                                24
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2; Page 12; 39pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Xie Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BIOWINDOW
                                                                                           SR,
                                                                                                                                                                                                                                                                                                                                                                                                            TCACTGCAACCTCTGCCTCCCGG 697
                                                                                                                                                                                                                                                                                                                                           standard;
                                                                                                                                                                                                                                                                                                                                                                                                TCACTGCAACCTCTGCCTCCCAG 2
                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                   (first
                                                                                             Bloch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENE
                                                                                                                                                                                                                                                                                                                                          DNA;
                                                                                                                                                                                                                                                        microsatellite amplification; genetic analysis; PCR;
                                                                                             ξ
                                                                                                                                                                                                                                                                                                   entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                    2.2%;
95.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3 C; 11 G; 5 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DEV
                                                                                                                                                                                                                                                                                                                                             24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INC
                                                                                                                                                                                                                                                                                                                                                                                                                                         0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 21.4;
Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SHANGHAI
                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                     1.1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length
                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                24;
                                 the
                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
                                          genetic
with an
                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             used
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Disclosure;

Page

29;

60pp; English

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1005 CGATTCTCCTGTCTCAGCCTCCC 1027

Matches Query Match

Local

Similarity

2.2%;

Conservative

0

Mismatches

DB 1;

Length 24;

Indels

0

Gaps

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RESULT 388
ADL06343/c
ID ADL063
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local S
Matches 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention relates to a method of reducing stutter in the amplification of a microsatellite. The method involves providing a sample comprising a microsatellite of interest; contacting the sample with at least one enzyme having nucleic acid polymerase activity and incubating the sample with the enzyme for amplifying the microsatellite. The method is useful for reducing stutter in the amplification of a microsatellite for genetic analysis. The present sequence is a PCR primer used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                               containing a site-specific recombinase characteristic sequence fragment, and the polynucleotide sequence encoding it. Also disclosed is a process for preparing the polypeptide by a DNA recombination technique and application of the polypeptide and polynucleotide in treating diseases such as growth development disorders and tumours. The present sequence represents a reverse transcriptase-PCR primer used in the examples of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; protein-13.2; site-specific recombinase;
growth.development disorder; tumour; reverse transcriptase-PCR;
                                                                                                                                                                                                                                                                                                                                                                                                            29-JAN-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                      CN1393548-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RT-PCR primer #1 for cDNA encoding human protein-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADL06343;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADL06343
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  illustrate the method of the invention
                                                                                                                                                                                              Example 3; SEQ ID NO 3; 32pp; Chinese.
                                                                                                                                                                                                                        Polypeptide-human protein-13.2 containing site specific recombinase characteristic sequence fragment and polynucleotide for coding it.
                                                                                                                                                                                                                                                                                               Mao
                                                                                                                                                                                                                                                                                                                                                   29-JUN-2001; 2001CN-00113178.
                                                                                                                                                                                                                                                                                                                                                                             29-JUN-2001; 2001CN-00113178
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                           Sequence 24
                                                                                                                                                                 The present invention relates to the isolation of human protein-13.2
                                                                                                                                                                                                                                                                                                                         (BIOW-)
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                                                                                                                                                                                                                                                                     2003-422181/40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                Xie
                                                                                                                                                                                                                                                                                                                         BIOWINDOW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AATCTCGGCTCACTGCAACCTCT 988
                                                                       invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24 BP; 7 A; 3 C; 9 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AATCTCGGCTCACTGTAACCTCT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                           BP; 6 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first
                                                                                                                                                                                                                                                                                                                          GENE DEV INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2.2%;
                                            4 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       24
                                            12 G; 2 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP.
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Pred.
Score 21.4; Db 1,
                                                                                                                                                                                                                                                                                                                            SHANGHAI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21.4;
No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               .1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ВB
                                            0 Other,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              24;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       a sample
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CGATTCTCCTGCCTCAGCCTCCC 2

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Query Match
Best Local Similarity
Matches 22; Conserv
                                                                                                                                                                                                                                                                                                                                                         Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide primer extension (SNPE) primers, and the sequences of regions flanking sites of single nucleotide polymorphisms SNPs. The present invention includes kits for determining the presence or absence of a SNP, using the oligonucleotides of the invention. The PCR primers are used to amplify a SNP flanking sequence, the SNPE primer is used as a genotyping primer. The oligonucleotides are useful for genotyping a nucleic acid sample by performing a single-nucleotide primer extension reaction. The oligonucleotides are useful for determining the presence, absence or identity of a SNP and for genotyping nucleic acid samples, for e.g. to assess by association analysis the genotype of an individual or group of individuals, having a pathological phenotype trait suspected of being caused by one or more SNPs. Phenotypic traits include diseases e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis;
                                                                                                                                             inflammation, cancer, nervous system diseases and infection by pathogenic microorganism. The method is also useful in forensic investigations and paternity analysis. The present sequence represents a single nucleotide primer extension (SNPE) primer specific for a human SNP containing DNA
                                                                                                                                                                                                                               agammaglobulinaemia, diabetes insipidus, Lesch-Nyhan syndrome, muscul dystrophy, familial hypercholesterolaemia, polycystic kidney disease, osteogenesis imperfecta and the intermit, polycystic kidney disease, osteogenesis imperfecta of controlation in the component of or susceptibility to multifactorial disease of which a component is or may be genetic such as autoimmune diseases, including, rheumatoid arthritis, multiple sclerosis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; Page 57; 83pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2001-290930/30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200129262-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SNP specific SNPE primer SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAH38671;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15-OCT-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        inflammation; forensic investigation; paternity analysis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (ORCH-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            sample.
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    Conservative
                                                                                      BP; 7
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                                                                                    Α,
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                      2.2%;
                                                                                    1 C; 5 G; 12 T; 0 U; 0 Other;
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                   Score 21.4; DB 1
Pred. No. 1.1e+03
    Mismatches
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                                          DB 1; Length 25;
    Indels
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                                                                                                                                                                                                                                                                                                                                              muscular
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Gaps

0

Query Match Best Local Similarity

2.2%; 1 C; 5

Score 21.4; DB 1; Pred. No. 1.1e+03;

Length

Sequence

25

7 Ą

G; 12 T; 0 U;

0 Other;

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CC primer extension (SNPE) primers, and the sequences of regions flanking concludes kits for determining the presence or absence of a SNP, using the presence or absence of a SNP, using the coligonucleotides of the invention. The PCR primers are used to amplify a SNP flanking sequence, the SNPE primer is used as a genotyping primer. CC The oligonucleotides are useful for genotyping a nucleic acid sample by performing a single-nucleotide primer extension reaction. The CC oligonucleotides are useful for determining the presence, absence or clidentity of a SNP and for genotyping nucleic acid sample by performing a pathological phenotypic trait suspected of being CC assess by association analysis the genotypic trait suspected of being CC agammaglobulinaemia, diabetes insipidus, Lesch-Nyhan syndrome, muscular cystrophy, familial hypercholesterolaemia, polycystic kidney disease, osteogenesis imperfecta and acute intermittent porphyria. Phenotypic traits also include symptoms of or susceptibility to multifactorial consists and including, rheumatoid arthritis, multiple sclerosis, inflammation, cancer, nervous system diseases and infection by pathogenic microorganism. The method is also useful in forensic investigations and commerce extension (SNPE) primer specific for a human SNP containing DNA commerce extension (SNPE) primer specific for a human SNP containing DNA commerce extension (SNPE) primer specific for a human SNP containing DNA commerce.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      밁
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequences AAH37205 - AAH40944 represent PCR primers,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; Page 55; 83pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     acid sample.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Picoult-Newburg L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis; inflammation; forensic investigation; paternity analysis; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SNP specific SNPE primer SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAH38231 standard; DNA; 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2001-290930/30.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              13-OCT-2000; 2000WO-US028436.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           14-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (ORCH-) ORCHID BIOSCIENCES INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              769
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TTTTTGTATTTTTAGTAGAGATG 791
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SNP; single nucleotide primer extension; naemia; diabetes insipidus; cancer;
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RESULT 391
AAZ45143/c
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                                                                                             implicated in the pathology of multiple sclerosis. Certain polymorphic sequences in the MMP-9 promoter, coding sequence and 3' untranslated region of the human MMP-9 gene (see AAZ45145) can affect the severity of atherosclerosis. The invention relates to the presence or absence of one variant form of a MMP-9 gene polymorphism (-1562 Cytosine/Threonine), detection of this polymorphism using oligonucleotides AAZ45137-Z45140 can be used for disease prognosis. The invention shows that the MMP-9 C-1562T polymorphism is a regulatory functional polymorphism. The methods and oligonucleotides are used to detect polymorphism in the MMP-9 gene. They are useful for the diagnosis and prognosis of diseases characterized by metalloproteinase mediated remodelling, such as atherosclerosis, tumour invasion and metastasis, inflammatory disease, and neurological diseases, particularly those involving demyelination such as multiple sclerosis, and afformation the MMP-9 gene variants may be and archiritic disease. Proteins encoded by the MMP-9 gene variants may be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matrix metalloproteinase-9; MMP-9; polymorphism; endopeptidase; detect; inflammatory disease; diagnose; atherosclerosis; tumour; metastasis; neurological disease; multiple sclerosis; arthritis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAZ45143 standard; DNA; 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     11-NOV-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Oligonucleotide used
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Detection of prognosis of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   07-MAY-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO9957315-A2
                                                                                                                                                                                                                                                                                                                         MMP-9 is a zinc-dependent endopeptidase, and is located on chromosome MMP activity is associated with inflammatory diseases and MMP-9 is implicated in the pathology of maintain.
                                                                                                                                                                                                                                                                                                                                                                          Oligonucleotides AAZ45143-Z45144 are used to determine the function the matrix metalloproteinase-9 (MMP-9) gene -1562 (C/T) polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                            Example 3; Page 14; 29pp; English
Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (ISIS-) ISIS INNOVATION LTD
                                                   ed for screening compounds that bind specifically to a molecule encoded one variant of a polymorphic sequence, thus identifying compounds ich modulate the activity of the enzyme. Such compounds can then be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  769
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP,
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   26
                                   rational drug
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             matrix metalloprotease diseases characterized
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   Α.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         to determine the function of MMP-9 polymorphism.
 7 C; 9
                                    design
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   င္ပ
   5 T;
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   0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             9 gene polymorphisms for diagnosis
by metalloproteinase mediated
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     0 Other;
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Query Match Best Local S Matches 23

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Gaps

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DB 1; Length 26;

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        cancers (e.g. adenocarcinoma, lymphoma, prostate cancer or uterus cancer), immune response, graft-versus-host disease, acquired immunodeficiency syndrome (AIDS), asthma, Crohn's disease, hypertension, congenital heart defects, multiple sclerosis, inflammation or Albright hereditary osteodystrophy and many other diseases listed in the specification. The DNA encoding the protein is useful in gene therapy for treating the conditions. This is also useful in detection assays, or chromosome mapping, tissue typing, diagnostic or prognostic assays, or for developing a powerful assay system for functional analysis of various human disorders, as well as in diagnostic applications. The present sequence is a primer used to isolate DNA encoding a NOVX protein by the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       encoding NOVX (or its complement, fragment or variant). NOVX is NOV1-14, 15a, 15b, 16a, and 16b. The NOVX polypeptide, nucleic acid encoding it and antibody against it, are useful for treating or preventing (e.g. by gene therapy) a NOVX-associated disorder in humans, e.g. cardiomyopathy, atherosclerosis, a disorder related to cell signal processing and metabolic pathway modulation, diabetes or cancers. The NoVX polypeptide and nucleic acids are also useful for determining the presence of predisposition to the diseases. The NoVX nucleic acid and polypeptide are especially useful in therapeutic or prophylactic applications for disorders associated with aberrant NoVX expression or activity, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; ss; NOVX; gene therapy; cardiomyopathy; atherosclerosis; cell signal processing disorder; metabolic pathway modulation disorder; diabetes; cancer; adenocarcinoma; lymphoma; prostate cancer; primer; uterus cancer; immune response; graft-versus-host disease; Exon linking; acquired immunodeficiency syndrome; AIDS; asthma; Crohn's disease; hypertension; congenital heart defects; multiple sclerosis; inflammation; hypertension; congenital heart defects; multiple sclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25-AUG-2000; 2000US-0228191P-
08-FEB-2001; 2001US-0267300P-
20-FEB-2001; 2001US-0269961P-
20-MAR-2001; 2001US-0277337P-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human NOV3
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                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to an isolated polypeptide (NOVX) a mature form of NOVX, a NOVX variant (differing by no more than 15%), the nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 1; Page 204; 263pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New polypeptides for treating or preventing a disorder associated with them, in humans, e.g. cardiomyopathy, atherosclerosis or cancers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Spytek
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               27-AUG-2001; 2001WO-US026510
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Albright hereditary osteodystrophy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Tomlinson JE,
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                                               cc acids (NAs) comprising contacting a NA sample from a physiological csource, with a pool of 50 distinct gene specific primers under suitable conditions to enzymatically generate sub-population of NAs, where each ceach labeled NA is generated using a single gene specific primer. The care method is useful for producing a sub-population of labeled NAs which is cuseful for analysing the differences in the RNA profiles between several cd different physiological sources, where the method comprises producing cuseful for analysing the differences in the RNA profiles between several cd different physiological sources, where the method comprises producing comprising the populations for each physiological sources, comprising the population, where the comparison is preferably comprising the population, where the comparison is preferably comprising the population, where the comparison is preferably comprised by hybridising the labeled NAs for each of the distinct comparison of a substrate to produce a hybridisation pattern for each of the sources, and comparing the patterns for each of the sources, where cd differential gene expression assays are utilised in differential content tissue or subtissue types. The present sequence is a threat of the source data for this patent did not form part of the printed comparing the pattern did not form part of the printed content the sequence data for this patent did not form part of the printed uspectification, but was obtained in electronic format directly from USPTO are betteronic format as a content part of the printed uspectification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to producing a sub-population of labeled nucleic acids (NAs) comprising contacting a NA sample from a physiological
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Producing sub-population of labeled nucleic acids, useful for analyzing differences in RNA profiles between several different physiological sources, using set of distinct gene specific primers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-314699/35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Chenchik A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Primer; ss; DNA microarray; differential expression analysis; human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human gene specific PCR primer #1216
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (CLON-) CLONTECH LAB INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local
                                    http.wipo.seqdata.uspto.gov/sequence.html?DocID=6352829B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                535 CTCCTGCCTCAGCCTCCCAAGTAGCT 560
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ID NO 1216; 11pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ВÞ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 21.2;
Pred. No. 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 26;
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Sequence

26

BP; 7

A; 5 C; 9 G; 5 T; 0 U;

0 Other;

21.2;

B

Length 26;

밁 Ś

26

Matches

Local Similarity es 23; Conserv

Conservative

0

Mismatches

Indels

0

Gaps

0

1.2e+03

867 GGGATTACAGGCGTGAGCCACCACGC 892

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RESULT 394
ABZ22656/c
ID ABZ226
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                              Query Match
                                                                                                                                                                                                                                                                                                            The present invention describes a method (M1) of screening for agents, conjugates or conjugate moieties (I), for transport by PEPT2 (an intestinal peptide transporter) transporter (II), comprising poviding a cell expressing (II), contacting the cell with (I), and determining if (I) passes into and/or through the cell by the way of (II). Also described: (I) a conjugate (III), comprising an agent linked to a conjugate moiety that is a substrate for (II), where the conjugate shows a Vmax of at least 1 % of Gly-Sar for (II), where the agent has a pharmaceutical activity without the conjugate moiety, and the conjugate has a greater Vmax for PEPT2 than the agent without the conjugate moiety; and conjugate moiety; and conjugate moiety and conjugate moiety; and conjugate moiety.
                                                                                                                        agent to a conjugate moiety to form a conjugate, where the conjugate is transported by (II) with a Vmax of at least 1 % of the Vmax of the substrate Gly-Sar, and formulating the conjugate with a carrier as a pharmaceutical composition. (III) is useful for treatment, by orally administering (III) to a patient, where the agent exerts a pharmacological effect in the patient who is free of a disease of brain, kidney, lung or spleen. The present sequence represents a PCR primer for human PEPTI, which is used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Screening for agents, conjugates or their moieties for transport by PEPT2 transporter, by contacting cell expressing transporter with the agent, and detecting their passage pass into and/or through the transporter.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      11-JUN-2001;
01-MAR-2002;
                                                                             Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Zerangue N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             31-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABZ22656 standard; DNA; 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example; Page 23; 43pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               11-JUN-2002; 2002WO-US018686
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; PEPT1; PEPT2; intestinal peptide transporter; transport;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human PEPT1 PCR primer PEPT1#1 R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABZ22656;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (XENO-) XENOPORT INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               867
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GGGATTACAGGCGTGAGCCACCACGC 892
                                                                             26 BP; 5 A; 9 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Dias T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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2002US-0361002P
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2.1%;
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       Score 21.2;
Pred. No. 1
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); Mismatches 3
                                DB 1;
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                              Length
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RESULT 395
ADI12547/c
                                                         RESULT 396
AAT62348/c
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                                                                                                                                                                                                                                                                    This invention relates to a novel method for predicting a predisposition to cancer in a patient by detecting large deletions in the human tumour converges of the pressor gene identified as the breast cancer susceptibility gene 1 (BRCAI). Specifically, it refers to deletions that result from the cumequal crossover between a pair of repetitive Alu sequences in the BRCAI consisted the recombined nucleotide sequence containing the confidence of the predisposition to breast and ovarian cancer. The present invention describes newly discovered deletion mutations that are consisted to be deleterious and cause significant alterations in the structure or biochemical function of BRCAI. Accordingly, it provides methods for detecting such mutants, as well as identifying and screening for cytostatic compounds useful for treating or preventing cancers associated with a BRCAI genetic variant. This polynucleotide is a mutant human BRCAI genomic DNA fragment that arises as a result of a crecombination event (deletion 4), which causes the omission of exons 16 cand 17, given in an exemplification of the invention.
                                                                                                                                                                                                 Query Match
Best Local
                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    breast cancer susceptibility gene 1; BRCA1; repetitive ovarian cancer; recombination; mutant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mutant human BRCA1 genomic DNA resulting from deletion 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Predicting a predisposition to cancer in a patient comprising detecting deletion in the BRCA1 gene that results from the unequal crossover between a pair of repetitive sequences in the BRCA1 gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               07-JUN-2002; 2002US-0387132P.
09-AUG-2002; 2002US-0402430P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ds; cancer; human; tumour suppressor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22-APR-2004
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                AAT62348
                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure;
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                                              AAT62348
                                                                                                                                                                                 Local Similarity
nes 23; Conserv
                                                                                                                                                     665
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                                                                                                                                                     CAATCTTGGCTCACTGCAACCTCTGC 690
                                              standard; DNA; 21 BP
                                                                                                                                                                                                                                               BP; 7 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQ ID NO 30; 59pp; English.
                                                                                                                                                                                  Conservative
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                                                                                                                                                                                          2.1%;
                                                                                                                                                                                                                                              6 C; 8 G; 5 T; 0 U; 0 Other;
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                                                                                                                                                                                                 Score 21.2;
Pred. No. 1
                                                                                                                                                                                  Mismatches
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                                                                                                                                                                                                             Length 26;
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RESULT 397
AAF95738
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to the amplification of region of DNA containing interspersed repetitive elements (IRE) such as the Alu repeat sequence (AAT62346). The method involves ligating a double stranded DNA structure with a non-complementary region, a 'bubble', in the centre (e.g. see AAT62343-4), to restriction digested fragments of regions containing IREs. The ligation results in a double stranded DNA molecule containing at least one 'bubble' at either end. After denaturing the structure, amplification of the IRE-containing region proceeds by PCR using primers targeted to the IRE sequence (e.g. AAT62347-50) and to the sequence in the 'bubble' region (e.g. see AAT62345). The primer presented here binds to nucleotides 216-236 of the Alu-S polymorphic repeat sequence. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bubble; interspersed repetitive element; ligation; annealing; primer; PCR; polymerase chain reaction; amplification; chromosomal aberration; genetic disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             using
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Amplification of nucleic acid having interspersed repetitive element -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     07-OCT-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US5597694-A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 21 BP; 5 A; 3 C; 9 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           method can be used to detect the presence or absence of a chromosomal aberration e.g. in a genetic disorder, in a test organism
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Col 17; 16pp; English.
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                                                                           Human; variant thrombospondin 1; variant thrombospondin 4; SNP; polymorphism; vascular disease; coronary artery disease; forensics; myocardial infarction; atherosclerosis; stroke; venous thromboembolism; pulmonary embolism; paternity test; ds.
                                                                                                                                                                                                                AAF95738;
                                                                                                                                                                                                                                              AAF95738 standard; DNA; 21 BP
 Variation
                                              Homo
                                                                                                                                                 Human gene single nucleotide polymorphism #499
                                                                                                                                                                                 06-JUN-2001
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                                              sapiens
                                                                                                                                                                                                                                                                                                                                                      967 ATCTCGGCTCACTGCAACCTC 987
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                bubble oligo:nucleotide
                                                                                                                                                                                                                                                                                                                          21 ATCTCGGCTCACTGCAACCTC
                                                                                                                                                                                                                                                                                                                                                                                      21;
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                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
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                                                                                                                                                                                   (first entry)
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Location/Qualifiers replace(11,C)
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                                                                                                                                                                                                                                                                                                                                                                                                                     2.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                     Score 21;
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; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1;
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RESULT 398
AAH24567/c
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Best Local Similarity
Matches 21; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention provides a method of diagnosing a vascular disease in an individual, involving determining the sequence at various polymorphic sites within the human thrombospondin 1 and thrombospondin 4 genes. The sequences at a number of polymorphic sites are also provided in the specification. In particular, the method can be used in the diagnosis of atherosclerosis, myocardial infarction, coronary heart disease, stroke, peripheral vascular diseases, venous thrombosembolism and pulmonary embolism. Single nucleotide polymorphisms (SNPs) are also useful in forensics, paternity testing, genetic analysis and phenotype correlations to diseases. The present sequence is an example of one of the human gene SNPS shown in the specification
04-AUG-1999;
                        04-AUG-1999;
                                                                                                                                                                                                                07-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           atherosclerosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nucleic acids comprising single nucleotide polymorphisms, useful applications such as forensics, paternity testing, medicine, generallysis and phenotype correlations to diseases such as diabetes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (WHED )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10-SEP-1999; 99US-0153357P.
26-JUL-2000; 2000US-0220947P.
16-AUG-2000; 2000US-0225724P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         07-SEP-2000; 2000WO-US024503.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO200118250-A2
                                                   08-MAY-2001.
                                                                                                         Homo
                                                                                                                                                         Human; Alu; metastatic potential determination; cancer;
                                                                                                                                                                                     Human Alu
                                                                                                                                                                                                                                                                    AAH24567 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example; Page 83; 242pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15-MAR-2001.
                                                                                                                                              choricallantoic
                                                                             US6228345-B1
                                                                                                      sapiens
                                                                                                                                migration; drug screening;
                                                                                                                                                                                                                                                                                                                                                             383
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WHITEHEAD INST BIOMEDICAL MILLENNIUM PHARM INC.
                                                                                                                                                                                                                                                                                                                                    CCTCCCAAAGTGCTGGGATTA 21
                                                                                                                                                                                                                                                                                                                                                            CCTCCCAAAGTGCTGGGATTA 403
                                                                                                                                                                                     sequence-specific primer Alu-Sense.
                                                                                                                                                                                                                                                                                                                                                                                                                                           BP; 5 A; 6 C; 5 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gargill M,
                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                              (first
99US-00366840
                        99US-00366840
                                                                                                                                              membrane; CAM; avian embryo;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /*tag= a
/standard_name= "single nucleotide polymorphism"
                                                                                                                                                                                                                entry)
                                                                                                                                                                                                                                                                                                                                                                                                    2.1%;
                                                                                                                                                                                                                                                                    21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ireland
                                                                                                                                                                                                                                                                    ₽₽
                                                                                                                                                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                                                                                                                                                                                                                    Score 21;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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                                                                                                                                PCR primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bolk S,
                                                                                                                                                                                                                                                                                                                                                                                                    DB 1; Length 21; 1e+03;
                                                                                                                                              intravasation;
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                                                                                                                                                                                                                                                                                                                                                                                       Indels
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(MOUN ) MOUNT SINAI SCHOOL MEDICINE
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Determining the metastatic potential of cancer cells and measuring invasion, comprises introducing cancer cells into the upper chorioallantoic membrane (CAM) and detecting cancer cell migration the upper CAM to the lower CAM. from

WPI; 2001-342659/36

Example; Col 11; 24pp; English.

comprises introducing a canner cell sample into the upper chorioallantoic membrane (CAM) of an avian embryo into which an artificially generated cair pocket has been created, incubating the embryo for intravasation to occur, and detecting migration of the cancer cells from the upper CAM to the lower CAM. The present sequence was used to selectively amplify human specific Alu repeat sequences, which will be present in the cancer cell CAM. DAM but not in the DNA of the CAM. This procedure enables detection of is useful for measuring the metastatic potential of cancer cells, for measuring the ability of the cancer cells into the lower CAM. The method cancer cells to invade blood vessels, and as a drug screening assay for the identification of agents having anticancer cells. The method may also be used to screen for agents capable of cancer cells. The method may also be used to screen for agents capable of cinhibiting cancer cell intravasation, and to detect phenotypic changes. CC effected by genetic manipulation of cancer cells that result in changes The invention relates to a method potential of cancer cells derived for determining the metastatic from a subject with cancer. The with cancer. The method

Sequence 21 BP; 5 P. 8 C; w ຸດ 5 T; 0 U; 0 Other;

Matches Query Match Local 21; Similarity Conservative 2.1%; 100.0%; 0, Score 21; Score 21; DB 1; Length 21; Pred. No. 1e+03; Mismatches 0 Indels 0; Gaps 0,

밁 390 AAGTGCTGGGATTACAGGCGT 410 21

S

RESULT 399 ABS98163

ABS98163 standard; DNA; 21 ΒP

ABS98163;

23-DEC-2002

(first entry

Human multidrug resistance gene polymorphic sequence #65

cyclooxgenase 2; COX2; diazepam binding inhibitor; DBI; haematological; epoxide hydroxylase 2; EPHX2; 5-lipoxygenase activating protein; FLAP; glutathione-S-transferase; EPHX2; 5-lipoxygenase activating protein; FLAP; glutathione-S-transferase; 12; GST12; histamine-N-methyl transferase; NNWT; MADPH quinone oxidoreductase 2; NQO2; sulfotransferase thermolabile; STM; UDP-glucuronosyl transferase 2B4; UDP-glucuronosyl transferase 2B7; UGT2B7; UDP-glucuronosyl transferase; UGT2B7; UDP-glucuronosyl transferase; UGT2B7; upp-glucuronosyl transferase; UGT2B7; upp-glucuronosyl transferase; UGT2B7; upp-glucuronosyl transferase; UGT2B7; upp-glucuronosyl transferase; up Human; ds; cytochrome P450 A1; CYP4501A1; UGT2B4; MDR1; cytochrome P450 A2; CYP4501A2; cytochrome P450 02E; CYP45002E1; LTF; adrenergic receptor beta1; ADBR1; aryl hydrocarbon; AHR; MRP3; NR112; aryl hydrocarbon receptor nuclear translocator; ARNT; cathepsin S; CTSS;

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X EX SOND NO SON
                                                                                                                                                                                                                                                                                                                                                                                                                                              cc identifying the genes responsible for a variety of disorder-related CC traits as a result of their e.g., overexpression, constitutive capression, mutation or underexpression, which may be used in diagnosing cc and/or treating the disorders. The nucleic acid molecules comprising the CC and/or treating the disorders. The nucleic acid molecules comprising the CC annur permanence of the disorders. The nucleic acid molecules comprising the CC annur permane and for MDR3 are useful for screening individuals for altered drug CC metabolism. The polymorphic sequences contained in CYP4501A1, CYPP4501A2, CC AHR, MDR1 and/or MDR3 may also be used to screen individuals for CC ausceptibility to cancer. Polymorphic sequences in ADRB1 or CHMR2 are CC used to screen for altered cardiovascular function, in COXZ for altered central corrous system function, in FLAP and HAWT for altered serine CC immunological or haematological function, in KLKZ for altered serine CC protease activity in the prostate, in LTF for altered immunological or contains and contains an
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                                                                                                                                                                                                                  Query Match
Best Local
                                                                                                                                                                         Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO200257410-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               transferase (UGT2B15), urokinase receptor (uPA), multidrug resistance 1 (MDR1), lactotransferrin (LTF), multidrug resistance associated protein 3 (MRP3), orphan nuclear receptor (NR112), or acetylcholine muscarinic receptor 1, 2, 3, 4, or 5 (CHMR1, CHMR2, CHMR3, CHMR3, CHMR4 or CHMR5) sequence. The polymorphisms in the human genes cited in the invention are useful as genetic linkage markers for locating and characterising the genes that are responsible for specific traits within the genome and eventually
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             28-NOV-2000; 2000US-00724389
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      protein (FLAP), glutathione-S-transferase 12 (GST12), histamine-N-meth
transferase (HMMT), (kallikrein 2) KLKZ, nicotinamide -N-methyl
transferase (NMMT), NADPH quinone oxidoreductase 2 (NQO2),
sulfotransferase thermolabile (STM), UDP-glucuronosyl transferase 2B4
(UGT2B4), UDP-glucuronosyl transferase 2B7 (UGT2B7), UDP-glucuronosyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Isolated nucleic acid molecules having polymorphisms in known human genes e.g. cytochrome p450 and cathepsin S useful as genetic linkage markers for locating, identifying and characterizing the genes responsible for disorder-related traits.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-698522/75.
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                                                                                                                                                                                                                                                                                                                                         Sequence 21
                                                                                  868
                                                                                                                                                                    Similarity
21; Conser
                                                                                  GGATTACAGGCGTGAGCCACC 888
                                                                                                                                                                                                                                                                                                                                         BP; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                DNA sequence of the invention
                                                                                                                                                                2.1%;
larity 100.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                         Α;
                                                                                                                                                                                                                                                                                                                                         6 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                         0
                                                                                                                                                                                                              Score 21;
Pred. No.
    21
                                                                                                                                                                         Mismatches
                                                                                                                                                                                                              DB 1;
. 1e+03;
                                                                                                                                                                         0,
                                                                                                                                                                                                                                                       Length 21;
                                                                                                                                                                         Indels
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RESULT 400 ADF38789/c

10-JUN-2004

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RESULT 401
ADO55495
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to a novel polynucleotide which encodes a protein involved in tumour necrosis factor (TNF)-alpha induced apoptosis. The polynucleotide of the invention demonstrates antiapoptotic activity and may be useful during gene therapy as an antisense polynucleotide for suppressing the expression of the protein involved in TNF-alpha induced apoptosis and for elucidating the mechanism of TNF-alpha induced apoptosis. The current sequence is that of the human TNF-alpha induced apoptosis-related DNA of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel polynucleotide encoding protein involved in tumor necrosis factor induced apoptosis, useful as probe to acquire perfect length cDNA of generated to TNF-alpha induced apoptosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (GENO-) GENO FUNCTION KK.
(DOKU-) DOKURITSU GYOSEI HOJIN SANGYO GIJUTSU
(TAHI/) TAHIRA K.
(KAWA/) KAWASAKI H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-APR-2002; 2002JP-00098130
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antisense gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human TNF-alpha induced apoptosis-related DNA - SEQ ID 17.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1; SEQ ID NO 17; 18pp; Japanese.
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                                                                                                                                                                                                                                                                                HIV gene expression analysis primer SB704 following siRNA inhibition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADO55495 standard; DNA; 21 BP
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                                                                                                                                              Human immunodeficiency virus 1
                                                                                                                                                                                                                                                                                                                                                  26-AUG-2004
                                                                                                                                                                                                                                                                                                                                                                                                            ADO55495;
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                                                                                                                                                                                                   anti-HIV; virucide; gene therapy; small interfering RNA; genome; diagnosis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2.1%;
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human; ds.
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Pred. No.
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ADH133/C ID ADH133 AC ADH133 AC ADH133 AC ADH133 AC ADH133 AC ADH133 AC ADH16 AC ADH
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Best Local 9
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                           Disclosure;
                                                                         Predicting, diagnosing or prognosing malignant neoplasia by detecting least two markers, where the markers are genes from one or more chromosomal regions altered in malignant neoplasia,.
                                                                                                                                                                                                                                                                                                                                        21-MAY-2002; 2002EP-00010291.
13-FEB-2003; 2003EP-00003112.
                                                                                                                                                                                                                                                                                                                                                                                                                      09-MAY-2003; 2003EP-00010447.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26-NOV-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EP1365034-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            gastric cancer; colon cancer; oesophageal cancer; mesenchymal cance bladder cancer; non-small cell lung cancer; human; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          malignant neoplasia; cytostatic; breast cancer; ovarian cancer; gastric cancer; colon cancer; oesophageal cancer; mesenchymal cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human malignant neoplasia-related PCR primer SeqID244.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            11-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADH13395 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          interference (RNAi). The methods and compositions of the present invention are useful for the diagnosis, prevention and/or treatment of HIV infections. This sequence corresponds to a PCR primer to carry out
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           sequence complementary to a portion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; SEQ ID NO 18; 59pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22-NOV-2002;
04-FEB-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   24-NOV-2003;
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                                                                                                                                                                                                                                                                                       (FARB ) BAYER AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ew small interfering RNA (siRNA) comprising a sequence complementary to portion of the HIV genome to mediate RNA interference (RNAi), useful or diagnosing, preventing and/or treating HIV infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            the siRNAs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            time PCR to determine gene expression after expression interference he siRNAs of the invention. .
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                                                                                                                                                                                                                                   Munnes M,
                             SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 4 A; 3 C; 9 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2002US-0428631P.
2003US-0444893P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           relates to a small interfering RNA (siRNA) lementary to a portion of the HIV genome to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Jacque
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                                                                                                                                                                                                                                      Kallabis
                             244;
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                        267pp;
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Pred. No.
                                                                                                                                                                                                                                      Ξ
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                           English
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CC This invention relates to a novel method for the prediction, diagnosis, CC or prognosis of malignant neoplasia by the detection of at least two CC markers. The invention may also be useful for the development of CC cytostatic compounds through the regulation of the expression of a gene CC or activity of a protein associated with malignant neoplasia. The method CC is useful for prediction, diagnosis or prognosis of malignant neoplasia cuch as breast cancer, ovarian cancer, gastric cancer, colon cancer, CC osephageal cancer, mesenchymal cancer, bladder cancer or non-small cell Clung cancer. The polymucleotides and polypeptides defined in the CC specification, antisense polymucleotides targeting the polymucleotides, CC antibodies targeting either one of the polymucleotides or polypeptides, CC antibodies targeting malignant neoplasia. The disease trated is CC preventing or treating malignant neoplasia. The disease trated is CC preferably breast cancer. The present sequence is that of a PCR primer CC which was used in the exemplification of the invention.
Sequence 23
     BP; 6 A;
     3 C; 9 G; 4 T;
     0 U; 1 Other;
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Matches
                                                 Query Match
                                         Local
                667
23
                                l Similarity
21; Conserv
ATCTTGGCTCACTGCAACCTC
               ATCTTGGCTCACTGCAACCTC 687
                                 Conservative
                                         100.0%;
                                                  2.1%;
                                 0;
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Pred. No.
w
                                 Mismatches
                                          1.1e+03;
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                                                  Length
                                  Indels
                                                  23
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RESULT 403
AAV19046/c
ID AAV19046 standard;
                                                                                                                                                                                                     Alu
                                                                                                                                                                                                                                                          AAV19046;
09-JUL-1996;
                            09-JUL-1996;
                                                                                     WO9801573-A1
                                                                                                               Saccharomyces
                                                                                                                             Synthetic
                                                                                                                                                         circular yeast
                                                                                                                                                                                                                               28-JUL-1998
                                                                                                                                                                                                   PCR primer
                                                                                                                                                         primer; amplification; Alu repeat sequence; vector;
ular yeast artificial chromosome; YAC; ss.
                                                                                                                                                                                                                               (first entry)
                                                                                                               ф
 96WO-US011478
                          96WO-US011478
                                                                                                                                                                                                                                                                                        DNA;
                                                                                                                                                                                                                                                                                        24
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Sn (HSSn)

DEPT

HEALTH & HUMAN SERVICES

This is the nucleotide sequence for the PCR primer used in the amplification of the 3' fragment of the Alu repeat sequence, which is used as a probe in the method of the invention. It involves the creation and use of circular yeast artificial chromosome (YAC) to selectively clone specific nucleic acids from a background of mixed nucleic acids by introducing the vector(s) into B. coli cells. They can be used to rapidly isolate human DNA where only a part of the sequence of DNA is known. Using the methods large fragments of DNA can be easily cloned and

Example 2; Page 61; 117pp; English.

Preparation of yeast artificial chromosomes -

comprising yeast centromere, for recombination.

marker,

by in vivo recombination
cer, yeast telomere and

y vector

WPI; 1998-110234/10

Resnick

Æ,

Larionov

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Kouprina NY,

Perkins

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RESULT 404
AAA27181
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A novel method for simultaneously determining the level of a number of CC target polynucleotides in a sample has been disclosed. The method CC involves forming double stranded copies of the target sequence in direct proportion to the target levels in the original sample. The target CC sequence is copied using primer pairs designed to flank a defined region in the target sequence. The double stranded copies are then cleaved and CC reacted with either first or second adaptor sequences. The first and CC through the target sequences. The adaptor sequences are then removed to leave target sequences. The adaptor sequences are then removed to form ultimer allow expression levels to be determined. This method is useful for developing polynucleotide abundance level profiles for cells and CC tissues under various conditions, stages of development and and disease states, particularly where the target polynucleotide is present at low CC levels. The method may also be used in the discovery and evaluation of CC addition to the method described above, the invention also includes the polynucleotide and polypeptide of P2. P2 is thought to be a member of a convel chemokine family, denoted CXSC and may be associated with immune treatment of asthma, allergic rhinitis (hay fever), urticaria (hives), candidate there shock and conditions involving immune system

CC anaphylactic shock and conditions involving immune system

CC therapy. The human P2 gene has been localised to chromosome 5, within the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Measuring target polynucleotide sequences in biological samples by contacting sequence-selective primer pairs, forming conjugates with adaptor molecules, polymerizing target-identifier dimers and quantifying
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAA27181
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Reverse primer P2 for target sequence human P2 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11-SEP-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 99; 103pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2000-387825/33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Dolganov
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C chemokine; Chromosome 5q31; gene rhinitis; urticaria; anaphylactic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novikov
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91.7%;
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Pred.
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No. 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1;
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RESULT 406 AAF24627/c ID AAF246

AAF24627 standard; DNA; 24

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RESULT 405
AAI65251
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                                               Query Match
Best Local S
Matches 22
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cytokine gene cluster at 5q31. The present sequence is the reverse primer P2 for target sequence human P2 gene
                                                                                                                     The invention relates to an isolated polypeptide of human dihydroorotase is comprising a sequence of 137 amino acids or its fragment, analogue or derivative. The polypeptide and the polynucleotide encoding it are useful in the diagnosis and treatment of malignant neoplasm, haemopathy, human immunodeficiency virus (HIV) infection, immunological diseases and various inflammatory diseases. The present sequence is a primer used to isolate a polynucleotide encoding the polypeptide of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; dihydroorotase 15; cytostatic; virucidal; immunomodulatory; antiinflammatory; haemostatic; anti-HIV; gene therapy; cancer; haemopathy; human immunodeficiency virus; HIV; infection; immunological disease; inflammatory disease; PCR primer; ss.
                                                                                                                                                                                                                                                    New human dihydroorotase 15 for diagnosing and treating malignant neoplasm, hemopathy, human immunodeficiency virus infection, immunological diseases and various inflammations.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human dihydroorotase 15 PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     29-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                           26-MAR-2001; 2001WO-CN000439
                                                                                                                                                                                                                                                                                                                                                                                                                                    04-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAI65251 standard;
                                                                                                Sequence
                                                                                                                                                                                                                            Example 2;
                                                                                                                                                                                                                                                                                                       WPI; 2001-597115/67
                                                                                                                                                                                                                                                                                                                               Mao Y,
                                                                                                                                                                                                                                                                                                                                                                                27-MAR-2000; 2000CN-00115187
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                                                                                                                                                                                                                                                                                                                                                        (SHAN-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            636 TCTGTCACCCAGGCTGGAGTGCAG 659
                         926
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      22;
                                                                                                                                                                                                                                                                                                                                                          SHANGHAI BIOWINDOW GENE
                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TATGTCACCCAGGCTGGGGTGCAG
                   GGAATCTCACTCTGTTACCCAGGC 949
                                                                                                 24
 GGAGTCTCACTCTGTCACCCAGGC
                                                                                                                                                                                                                            Page 17; 36pp; Chinese.
                                                                                                 ВP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                  Conservative
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                                                                                                 4 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA;
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                                                          2.1%;
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                                                                                                   G; 5 T; 0 U;
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Pred. No. 1.
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                                                              Score 20.8;
Pred. No. 1
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RESULT 407
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Best Local Similarity
Matches 22; Conserv
             Homo sapiens
                                       3-hydroxy-3-methylglutaryl-coenzyme A reductase gene; dyslipidemia; HMG-CoA reductase gene; genetic marker; cardiovascular disease; myocardial infarction; stroke; PCR primer; ss.
                                                                                                                                                          AAF24635;
                                                                                                                                                                                                                                                                                                                                                                            Sequence 24 BP; 5 A; 8 C; 8 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                     useful for diagnosis of HMG-COA reductase mediated diseases such dyslipidemia and other cardiovascular diseases such as myocardial infarction and stroke. HMG-COA reductase antagonist drugs are use treat dyslipidemia and other cardiovascular diseases such as myoc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PCR primers AAF24627-28 were used to detect a polymorphism in the human 3 -hydroxy-3-methylglutaryl-coenzyme A (HMG-CoA) reductase gene. The polymorphism is present in the promoter region, exon 15, introns 2, 5, 15 or 18. HMG-CoA reductase polymorphisms are useful as genetic markers in linkage studies. Detection of the presence of the polymorphisms is useful for assessing the pharmacogenetics of therapeutic compounds in the treatment of HMG-CoA reductase mediated diseases. The polymorphisms are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 1; Page 31; 45pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel polymorphisms in human 3-hydroxy-3-methylglutaryl-coenzyme A (HMG-CoA) gene useful for diagnosis and treatment of HMG-CoA reductase-mediated diseases such as dyslipidemia and other cardiovascular diseases
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3-hydroxy-3-methylglutaryl-coenzyme A reductase gene; dyslipidemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Primer for a polymorphism at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20-APR-2001
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                                                                                                                             20-APR-2001
                                                                                                                                                                                     AAF24635 standard;
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                                                                                                                                                                                                                                                                                       931
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                                                                                                                                                                                                                                                                                       CTCACTCTGTTACCCAGGCTGGAG 954
                                                                                                polymorphism
                                                                                                                                                                                                                                                                                                                                                                                                          and stroke
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Pred. No. 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     marker; cardiovascular disease;
                                                                                                                                                                                                                                                           _
                                                                                                of human
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                                                                                               HMG-COA
                                                                                                                                                                                                                                                                                                                                               Length
                                                                                                                                                                                                                                                                                                                      Indels
                                                                                               reductase gene.
                                                                                                                                                                                                                                                                                                                                                 24;
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RESULT 408
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Matches
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                                                                                                                                                                                                                                                                                                                                                                       Human; reverse transcriptase 13; cytostatic; virucide; antiinflammatory; haemostatic; gene therapy; malignant haemopathy; HIV infection; immunological disease; infigure
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    dyslipidemia and other cardiovascular diseases such as myocardial infarction and stroke. HMG-CoA reductase antagonist drugs are use treat dyslipidemia and other cardiovascular diseases such as myoc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel polymorphisms in human 3-hydroxy-3-methylglutaryl-coenzyme A (HMG-COA) gene useful for diagnosis and treatment of HMG-COA reductase-mediated diseases such as dyslipidemia and other cardiovascular diseases
                        Mao Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human reverse
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAH75870;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAH75870 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
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                                                                                                               02-MAR-2000; 2000CN-00111806.
                                                                                                                                                             26-FEB-2001; 2001WO-CN000280
                                                                                                                                                                                                              07-SEP-2001
                                                                                                                                                                                                                                                         WO200164893-A1
                                                                                                                                                                                                                                                                                                                                                      developmental disorder; PCR primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   infarction and stroke
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                                                                      (BIOW-) BIOWINDOW GENE DEV INC SHANGHAI.
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                          Xie Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CTCACTCTGTGGCCCAGGCTGGAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   transcriptase 13 coding sequence PCR primer #2
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Pred. No. 1.2e
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                              inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                       immunomodulatory;
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RESULT 409
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Best Local S
Matches 22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to human reverse transcriptase 13 and its coding sequence (see AAH75868 and AAG66428). The reverse transcriptase and its coding sequence are useful in the diagnosis and treatment of malignant tumour, haemopathy, HIV infection, immunological diseases, various inflammations and developmental disorders. The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 3; Page 12; 34pp; Chinese
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       developmental
                          The invention relates to an isolated polypeptide of ribosome sl9e protein 11 and its corresponding coding sequence. The polypeptide and encoded polynucleotide are applicable in diagnosis and treatment of malignant tumours, haemopathy, human immunodeficiency virus (HIV) infection, immunological diseases and various inflammations. The present sequence represents the reverse transcriptase (RT) PCR primer #2 used in analysis of ribosome sl9e protein II
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; ribosome s19e protein II; cytostatic; virucidal; immunomodulatory; antiinflammatory; haemostatic; malignant tumour; haemopathy; ss; human immunodeficiency virus; HIV; immunological disease; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ribosome s19e
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAS12447
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                                                                                                                                                                diseases and inflammations.
                                                                                                                                                                            Ribosome s19e protein 11 and encoded polynucleotide for diagnosis and treatment of malignant tumors, hemopathy, HIV infection, immunologica
                                                                                                                                                                                                                       WPI; 2001-582156/65.
                                                                                                                                                                                                                                                                                                       02-MAR-2000; 2000CN-00111831.
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                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                               PCR primer.
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                                                                                                                                      Example
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                                                                                                                                                                                                                                                                              BIOWINDOW GENE
                                                                                                                                      2; Page 12; 34pp; Chinese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   protein 11, RT-PCR primer #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA;
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91.7%;
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Pred. No. 1.2e
0; Mismatches
                                                                                                                                                                                                                                                                              SHANGHAI.
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tumor, hemopathy, human
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                                                                                                                                                                                  immunological
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Other,

Human myosin 15-JAN-2002

heavy (first

chain 12-14 coding

sequence

PCR

primer

#1

entry)

AAI71673

standard; DNA;

24

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RESULT 410
AAI66532
ID AAI665
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RESULT 411
AAI71673/c
ID AAI716
XX
AC AAI716
XX
AC AAI716
XX
DT 15-JAN
XX
DE Human
XX
                                                                                                                      片
                                                                                                                                       S
                                                                                                                                                            Query Match
Best Local S
Matches 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
Matches 22; Conserv
                                                                                                                                                                                                                                 The present invention provides the protein and coding sequences of hum pterin-molybdenum oxidoreductase 10. The sequences can be used in the treatment of cancer, haemopathy, HIV infection, immunological diseases and inflammation. The present sequence is a PCR primer for the coding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAI66532 standard;
                                                                                                                                                                                                                                                                                                      New polypeptide for the diagnosis and treatment of malignant neoplasm, hemopathy, HIV infection, immunological diseases and inflammations, comprises the human pterin-molybdenum oxidoredctase 10 protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; pterin-molybdenum oxidoreductase 10; cancer; haemopathy;
immunological disease; HIV infection; inflammation; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human pterin-molybdenum oxidoreductase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAI66532;
                                                                                                                                                                                                                                                                                                                                                                   Mao
                                                                                                                                                                                                                                                                                                                                                                                                           24-MAR-2000; 2000CN-00115110
                                                                                                                                                                                                                                                                                                                                                                                                                               23-MAR-2001; 2001WO-CN000393
                                                                                                                                                                                                                                                                                                                                                                                                                                                   04-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200172788-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          11-DEC-2001
                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                         sequence
                                                                                                                                                                                                                                                                                   Example 2; Page 17; 36pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                WPI; 2001-602841/68
                                                                                                                                                                                                                                                                                                                                                                                       (SHAN-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                   ۲,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       181
                                                                                                                                          638
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ᆫ
                                                                                                                                                             22;
                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                   Xie
                                                                                                                                                                                                                                                                                                                                                                                      SHANGHAI BIOWINDOW GENE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TAGAGATGGAGTTTCTCCATGTTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TAGACATGGGGTTTCTCCATGTTG
                                                                                                                                                                                                    24
                                                                                                                                                                                                                         of the invention
                                                                                                                      TGTCATCCAGGCTGGAGTACAGTG
                                                                                                                                          TGTCACCCAGGCTGGAGTGCAGTG
                                                                                                                                                                                                    BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first
                                                                                                                                                                                                    5 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2.1%;
                                                                                                                                                                      2.1%;
                                                                                                                                                                                                    5 C; 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 20.8; Di
Pred. No. 1.2e
0; Mismatches
                                                                                                                                                                                                      ଦ
                                                                                                                                                              0
                                                                                                                                                          Pred. No. 1.25
0; Mismatches
                                                                                                                                                                         Score 20.8;
Pred. No. 1
                                                                                                                                                                                                      6 T;
                                                                                                                                                                                                                                                                                                                                                                                          DEV
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                                                                                                                        24
                                                                                                                                           661
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U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        10 cDNA PCR
                                                                                                                                                                       .2e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       .2e+03
                                                                                                                                                                                                       0 Other;
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                                                                                                                                                                                  ۲;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length
                                                                                                                                                                                 Length
                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   24
                                                                                                                                                                                     24;
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                                                                                                                                                                0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                    human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
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RESULT 412
AAF69722
ID AAF697
XX AAF697
XX AAF697
XX Human
XX Polymo
KW Polymo
KW Allerg
XX Homo 8
PN WO2001
XX Homo 18
PN W13-JUI
XX 13-JUI
XX 13-JUI
XX (GENA-
XX (GENA-
XX WPI; 2
PI Winden
XX WPI; 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          밁
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Best Local S
Matches 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo
  New isolated polynucleotide useful for the identification of therapeutics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention provides the protein and coding sequences of human myosin heavy chain 12-14. The sequences can be used in the treatment of prader Willi syndrome, Klinefelter syndrome, kinetic illnesses and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Peptide-human myosin heavy chain 12-14 and encoded polynucleotide, used in diagnosis and treatment of Prader Willi syndrome, and Klinefelters
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mao Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200185752-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; myosin heavy chain 12-14; Prader Willi syndrome; PCR primer; Klinefelter syndrome; inflammation; kinetic illness; gene therapy; ss.
                         WPI; 2001-103078/11.
                                                                                                                                                         18-JAN-2001
                                                                                                                                                                                 WO200104270-A1
                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                            allergic
                                                                                                                                                                                                                                                             Human IL4Ralpha gene PCR primer #58.
                                                                                                                                                                                                                                                                                       18-APR-2001
                                                                                                                                                                                                                                                                                                              AAF69722;
                                                                                                                                                                                                                                                                                                                                    AAF69722 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sequence of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 2; Page 17; 39pp; Chinese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              29-APR-2000; 2000CN-00115544.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28-APR-2001; 2001WO-CN000670
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15-NOV-2001
                                                  Windemuth
                                                                                                                                 13-JUL-2000; 2000WO-US019094
                                                                                                        13-JUL-1999;
                                                                                                                                                                                                                                         Polymorphism;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (SHAN-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2001-648982/74.
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                                                                                                                                                                                                                                                                                                                                                                                                                     969
                                                                                                                                                                                                                                                                                                                                                                                               24
                                                                                                                                                                                                                                                                                                                                                                                                                                            22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SHANGHAI BIOWINDOW GENE
                                                                                    GENAISSANCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          24
                                                 Denton RR, h AK;
                                                                                                                                                                                                                             disease;
                                                                                                                                                                                                                                                                                                                                                                                               CTCGGCTCACTGCAAGCTCCGCCT 1
                                                                                                                                                                                                                                                                                                                                                                                                                   CTCGGCTCACTGCAACCTCTGCCT 992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP; 5 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                       human; interleukin 4 receptor-alpha; IL4R-alpha;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present sequence
                                                                                                           99US-0143435P
                                                                                                                                                                                                                              PCR
                                                                                                                                                                                                                                                                                                                                     DNA;
                                                                                   PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          5 C; 11
                                                            Duda A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                        91.7%;
                                                                                                                                                                                                                             primer;
                                                                                                                                                                                                                                                                                                                                     24
                                                                                                                                                                                                                                                                                                                                     BP
                                                                                                                                                                                                                                                                                                                                                                                                                                            0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score
Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          <u>.</u>
                                                                                                                                                                                                                              88
                                                            Nandabalan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          w
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DEV
                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           syndrome,
                                                                                                                                                                                                                                                                                                                                                                                                                                                        20.8;
No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        .2e+03
                                                           ㅈ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             primer
                                                            Stephens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              for
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                                                                                                                                                                                                                                                                                                                                                                                                                                            <u>,,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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밁
                             S
                                                                    Matches
                                                                                                  Query Match
                                                                                                                                                                   The present invention relates to polymorphisms of the human interleukin 4 receptor-alpha gene (IL4R-alpha; see AAF57718 for the reference sequence). Polynucleotides comprising polymorphic gene variants are useful for therapeutic purposes. For example, where a patient may benefit from expression of a particular IL4Ralpha protein isoform, an expression vector encoding the isoform may be administered to the patient. It may desirable to decrease or block expression of a particular IL4Ralpha isogene, which may be done by turning off by transforming a targeted organ, tissue or cell population with an expression vector that expresses high levels of untranslatable mRNA for the isogene. Specific therapeutics identified by these methods may be useful for allergic diseases. The present sequence is a PCR primer for human IL4R-alpha
                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                           Example 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                               in allergic diseases is new
                  1002 AAGCGATTCTCCTGTCTCAGCCTC 1025
                                                                  l Similarity
22; Conser
Ľ
                                                                                                                                         24
AAGCGATTCTTCTGCCTCAGCCTC
                                                                                                                                                                                                                                                                                                                                                                                                         Page
                                                                                                                                         BP; 4 A;
                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                         62; 188pp; English.
                                                                                   2.1%;
91.7%;
                                                                                                                                         9 C; 4 G; 7 T; 0 U; 0 Other;
                                                                  0,
                                                                                   Score 20.8;
Pred. No. 1
                                                                    Mismatches
24
                                                                                   1.2e+03;
                                                                                                      DB 1;
                                                                      2
                                                                                                    Length
                                                                      Indels
                                                                                                        24
                                                                  0;
                                                                  Gaps
                                                                    0
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RESULT 413
AAI68386
                                                                                                                                                                                                                    03-JAN-2002
                                                                                                                                                                                                                                                                                     AA168386
11-OCT-2001.
                                 WO200175042-A2
                                                               Homo sapiens.
                                                                                                                immunomodulatory; antiinflammatory; haemostatic; malig
infection; human immunodeficiency virus; gene therapy;
                                                                                                                                                 Human; ATP-dependent hydrolase serine 9;
                                                                                                                                                                                   Human ATP-dependent hydrolase serine 9 PCR primer SEQ
                                                                                              immunological disease;
                                                                                                                                                                                                                                                                                       standard;
                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                       DNA;
                                                                                                  PCR
                                                                                                                                                                                                                                                                                       24
                                                                                                primer; ss.
                                                                                                                                                                                                                                                                                         ₽P
                                                                                                                                                cytostatic; virucidal;
                                                                                                                                   malignant
                                                                                                                                                                                     ij
                                                                                                                                                                                     ö
                                                                                                                                   tumour;
                                                                                                                                   HIV;
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26-MAR-2001; 2001WO-CN000434. 27-MAR-2000; 2000CN-00115164.

Mao

Xie Y;

(SHAN-)

SHANGHAI BIOWINDOW GENE

DEV

INC

WPI; 2001-626418/72.

The invention relates to human ATP-dependent serine hydrolase 9 with cytostatic, virucidal, immunomodulatory, antiinflammatory and haemostatic activity. The protein and encoding polynucleotide are used in diagnosis and treatment of malignant tumour, haemopathy, human immunodeficiency virus (HIV) infection, immunological diseases and various inflammations. The polynucleotide is useful in gene therapy. The present sequence is that of a PCR primer, useful to the invention

Human ATP-dependent serine hydrolase 9 and encoded polynucleotide, in diagnosis and treatment of malignant tumors, hemopathy, human immunodeficiency virus infection, immunological diseases and

Example 2; Page 17; 33pp; Chinese.

inflammation.

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RESULT 414
ABA82841/c
ID ABA828
XX ABA828
XX ABA828
XX ABA828
XX Human
XX Human;
KW Human;
KW neurol
KW ds.
XX O9-API
XX O9-API
XX U02001
XX WPI;
PR (COGE
XX WPI;
DR P-PSD;
X
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밁
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Best Local S
Matches 22
                                                                                               Matches
                                                                                                                                           Query Match
                                                                                                                                                                                                                                      The present invention relates to protective sequence proteins (ABB44624-ABB44830) and their coding sequences (ABA82701-ABB2937). The sequences, when introduced into a cell either predisposed to undergo cell death or in the process of undergoing cell death, prevent, delay or rescue the cell from death, hence, these sequences are named "protective sequences". The sequences are useful for treating and/or ameliorating cancer, autoimmune diseases and neurological disorders e.g. stroke. Further examples of diseases which may be treated by the present invention are given in the specification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABA82841 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New protective sequences and their products, useful for diagnosing treating diseases involving cell death, including neurological disease. stroke and for identifying modulators of expression of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 11-APR-2000; 2000US-00547735
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200176457-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; protective sequence; cell death; cancer; autoimmune disease; neurological disorder; stroke; cytostatic; neuroprotective; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          07-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABA82841;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  09-APR-2001; 2001WO-US011663
                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               protective sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           P-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2002-025874/03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      sapiens.
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                                                 177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 <u>ب</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    protective
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24
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В
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                                                                                                                      Similarity
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                                                                                                                                                                                               24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Fig 11;
                                                 TTAGTAGAGATGGAGTTTCTCCAT 200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP; 5 A;
                                                                                                                                                                                               BP; 7 A; 6 C; 4 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Portbury
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NEUROSCIENCE INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           283pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ე
8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SD,
                                                                                                                      2.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        5 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Puranam
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                                                                                               Score 20.8; I
Pred. No. 1.26
0; Mismatches
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Pred. No. 1.2e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Katz
                                                                                                                         .2e+03;
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                                                                                                                                                DB 1;
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                                                                                                                                              Length 24;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Barney
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                                                                                               Gaps
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RESULT 415 ABL59102/c

RESULT 416 ABK14186

ABK14186 standard; DNA;

24 В₽

of.

0

ABK14186;

21-MAY-2002

(first entry)

Human splicing

factor 9.24 cDNA RT-PCR primer

#2.

Human; splicing factor 9.24; ss; cytostatic; gene therapy; cancer; tumour; foetus deforming; protein metabolic disturbance related disease;

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                                                                         Query Match
Best Local S
                                                              Matches
                                                                                                                                                    method, designated transformation-associated recombination, eliminates the need for an in vitro ligation step, and makes possible selective cloning of cDNAs for which only the 3'-sequence is known. The method is used for making a YAC. The method is also used for selective cloning of mammalian, specifically human, nucleic acid from a population, particularly radiation hybrids that contain only a small fragment of a pluman chromosome. PCR primers ABL59102-03 were used to amplify an 82 bp Alu probe from the pPD39 plasmid containing an Alu consensus sequence. The probe was used to identify human YACs, generated using the method of the invention
                                                                                                                                                                                                                                                                                                              The specification describes a method for making a yeast artificial chromosome (YAC) that includes an origin of replication (ori). The metho comprises incorporating into yeast cells: a population of mammalian nucleic acid; and a vector that comprises a yeast centromere, selection marker, yeast telomere and a sequence that recombines with a region of the nucleic acid, so that in vivo recombination to a VAC occurs. This
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US6391642-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                transformation-associated recombination; PCR; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             27-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Preparing yeast artificial chromosomes, useful e.g. for cloning specific human nucleic acid, comprises recombination in yeast cells between a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    14-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21-MAY-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Yeast artificial chromosome; YAC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABL59102;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABL59102 standard; DNA; 24 BP
                                                                                                                       Sequence 24 BP; 4 A; 6 C; 11 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                            Example 2; Col 35; 50pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            nucleic acid and a yeast vector.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2002-498777/53.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Resnick MA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (USSH ) US
                   675 TCACTGCAACCTCTGCCTCCCGGG 698
 24 TCACTGCAAGCTCCGCCTCCCGGG 1
                                                            22;
                                                                           Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              used to amplify an 82 bp Alu probe
                                                            Conservative
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91.7%;
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                                                                            Score 20.8;
Pred. No. 1
                                                              Mismatches
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                                                                              2e+03
                                                                                             DB 1;
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                                                                                           Length 24;
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Best Local S
Matches 22
                                                   Mao
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           Human topoisomerase I
                               WPI; 2002-281738/33.
                                                                                                                                26-DEC-2001.
                                                                                                                                                   CN1328155-A
                                                                                                                                                                       Homo
                                                                                                                                                                                                 Human; topoisomerase I 9.79;
                                                                                                                                                                                                                     Human topoisomerase I 9.79 protein,
                                                                                                                                                                                                                                                                                 ABK12860
                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                 reverse transcription-PCR (RT-PCR) primer used in isolation of cDNA encoding the human splicing factor 9.24 polypeptide of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                             the DNA sequence encoding it. The DNA and protein sequences are undiagnosis and treatment of tumours, foetus deforming and protein metabolic disturbance related diseases. This sequence represents reverse transcription-PCR (RT-PCR) primer used in isolation of cl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 2; Page 12; 38pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human splicing factor 9.24 in diagnosis and treatment related disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19-JUN-2000; 2000CN-00116576
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14-FEB-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RT-PCR;
                                                                                        14-JUN-2000; 2000CN-00116475
                                                                                                           14-JUN-2000; 2000CN-00116475.
                                                                                                                                                                                         reverse transcriptase PCR.
                                                                                                                                                                                                                                           18-JUN-2002
                                                                                                                                                                                                                                                                ABK12860;
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                                                                     (BODE-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   invention relates to the human splicing factor 9.24 polypeptide
                                                                                                                                                                     sapiens
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                                                                     BODE GENE
                                                                                                                                                                                                                                                                                                                                                                                                                24
                                                                                                                                                                                                                                                                                  standard;
                                                                                                                                                                                                                                                                                                                                   AACTITGTCACCTAGGCTGGAGTG
                                                                                                                                                                                                                                                                                                                                                      AACTCTGTCACCCAGGCTGGAGTG
                                                                                                                                                                                                                                                                                                                                                                                                               BP; 5 A; 5 C; 7 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                          (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            transcription-PCR; primer.
teratogenesis
                                                                     DEV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENE DEV INC
                                                                                                                                                                                                                                                                                  DNA;
                                                                                                                                                                                                                                          entry)
                                                                                                                                                                                                                                                                                                                                                                                  2.1%;
91.7%;
           9.79
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                                                                     GLT
polypeptide and sais and tumors.
                                                                                                                                                                                                                                                                                  ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  polypeptide and encoding polynucleotide, used of tumors and protein metabolic disturbance
                                                                                                                                                                                                                                                                                                                                                                        0;
                                                                                                                                                                                                                                                                                                                                                                                 Score 20.8;
Pred. No. 1.
                                                                     SHANGHAI.
                                                                                                                                                                                                   teratogenesis;
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                                                                                                                                                                                                                                                                                                                                                                        Mismatches
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                                                                                                                                                                                                                       RT-PCR
                                                                                                                                                                                                                                                                                                                                                                                  .2e+03
                                                                                                                                                                                                                                                                                                                                                                                            DB 1;
           the
                                                                                                                                                                                                                      primer1.
                                                                                                                                                                                                   tumour;
          polynucleotide encoding
                                                                                                                                                                                                                                                                                                                                                                                            Length
                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                 primer;
                                                                                                                                                                                                                                                                                                                                                                                            24;
                                                                                                                                                                                                  ss; RT-PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          are used
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RESULT 418
ABZ25248/c
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Best Local S
Matches 22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 polypeptide, the polymucleotide encoding it and a DNA recombination process used to produce the polypeptide. The invention also discloses the agonist resisting the polypeptide. The polypeptide and its antagonist are useful for treating teratogenesis and tumours. The present nucleic acid sequence represents a reverse transcriptase (RT)-PCR primer that was used in the methods of the invention to isolate the coding sequence of the hyperbolic transcriptase the coding sequence of the primer that was used in the methods of the invention to isolate the coding sequence of the primer that was used to be a sequence of the invention to isolate the coding sequence of the primer that was used to be a sequence of the process of the invention that was used to be a sequence of the process of
                                                                                                                                                                    The present invention relates to human peroxidase 9.90 The peroxidase is useful for treating diseases such as infection. The present sequence is a PCR primer, which example from the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; peroxidase 9.90; enzyme; cancer; HIV infection; cytostatic; anti-HIV; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human peroxidase
                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                        Polypeptide-human peroxidase
                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-733654/80.
                                                                                                                                                                                                                                                                                                                                                                                                                             Mao
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20-DEC-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CN1360029-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  24-APR-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 2; Page 18 (Disclosure); 32pp; Chinese
                                                                                                                                                                                                                                                                             Example
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24-JUL-2002.
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    1099 CACCATATTTGTCAGGCTGGTCTC 1122
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                                             Similarity
22; Conserv
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                                                                                                                                                                                                                                                                             Page 16 (Disclosure); 31pp; Chinese
                                                                                                                              BP; 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    invention relates to a new human topoisomerase I 9.79
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                                             Conservative
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                                                                                                                            5 C; 8
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91.7%;
                                                               2.1%;
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                                             <u>,</u>
                                                                                                                          G; 4 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                        protein 9.90
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Pred. No. 1.2e+03;
                                                               Score 20.8;
Pred. No. 1
                                             Mismatches
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                                                               .2e+03
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                                                                                     DB 1;
                                                                                                                                                                                                                                                                                                                                          and polynucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 1;
                                                                                   Length
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                                                                                                                                                                                                               cancer
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cancer and HIV
                                                                                       24;
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                                             Gaps
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CACCATATTGCTCAGGCTGGTCTC

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RESULT 419
ABA02134
                                                                                                                                     CC unchein acids encoding it (ABA02133), and a method for the recombinant CC production of zinc ion transport protein 26. The protein has a molecular CC weight of 26 kD, and has 35% identity and 54% homology over a 210 amino CC acid stretch with the rat zinc transporter ZmT-1 (GenBank accession CC unmber U17133). The present invention additionally discloses an CC antipody which specifically binds to zinc ion transport protein 26. Zinc CC ion transport protein 26, and nucleotides which encode it may be used for CC treating a variety of diseases, such as malignant tumours, blood CC diseases, HIV (human immunodeficiency virus) infection, immune disorders, CC inflammatory conditions, embryonic development disorders, and development CC and growth disorders. The protein may also be used to screen for CC modulators of its activity or for peptide fingerprinting identification. CC modulators of its activity or for peptide fingerprinting identification CC reactions or as a probe for hybridisation reactions, or in producing gene chips or microarrays. Sequences ABA0213-ABA02135 represent reverse CC invention to isolate human sinc ion reactions, or in producing gene convention to isolate human sinc ion reactions of the human sinc ion reactions of the human sinc ion reactions of the invention to isolate human sinc ion reactions.
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Best Local
                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABA02134 standard; DNA; 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  immune disorder; inflammatory condition; embryonic development disorder; developmental disorder; growth disorder; cytostatic; anti-HIV; antiinflammatory; immunomodulator; reverse transcription-PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human zinc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  08-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human zinc ion transport protein 26 and encoded polynucleotide, diagnosis and treatment of malignant tumors, hemopathy, human immunodeficiency virus infection, immunological diseases and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        23-APR-2001; 2001WO-CN000610
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                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to human zinc ion transport protein 26 (AAM52621),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 3; Page 11; 31pp; Chinese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2002-026163/03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (BIOW-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    27-APR-2000; 2000CN-00115461
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      blood disease; HIV
                  1096 TTTCACCATATTTGTCAGGCTGGT 1119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      zinc ion transport protein 26; rat ZnT-1 homologue; ransporter; recombinant production; malignant tumour; disease; HIV infection; human immunodeficiency virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   primer;
                                                                l Similarity
22; Conserv
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                                                                                                                                     24
TTTCACCATATTGGCCAGGCTGGT 24
                                                                                                                                                                        to isolate human zinc ion transport protein
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                                                                2.1%;
larity 91.7%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     88.
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                                                                                                                                     A;
                                                                                                                                     6 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              protein 26 RT-PCR primer, SEQ ID
                                                                                                                                     δ
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                                                                  0;
                                                                                                                                     G; 8 T; 0 U;
                                                                Score 20.8; D
Pred. No. 1.2e
0; Mismatches
                                                                                                                                       0 Other;
                                                                                   2e+03;
                                                                                                      В
                                                                                                    1,
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                                                                                                                                                                        26 CDNA
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                                                                  Gaps
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RESULT 420

Human, plasminogen activator inhibitor 2-9.9; primer; 88; thrombosis; haemorrhagic disease; cerebral infarction; myocardial infarction; tum

tumour;

0

haemopathy; human immunodeficiency virus; HIV; inflammation;

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RESULT 421
ABS57470/C
ID ABS574
XX
ABS574
XX
ABS577
XX

Z7-FEE
DT 27-FEE
DT 27-FEE
DT Human
XX
Human
X
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to an isolated polypeptide of human microtubulin 11, the nucleic acid encoding it, a fragment, analogue or derivative of it, a transformed cell expressing the protein from an expression vector, antibodies against the protein and ant/agonists of the protein. The polypeptide and encoded polynucleotide are applicable in diagnosis and treatment of cancer, haemopathy, human immunodeficiency virus infection, treatment of cancer, haemopathy, human immunodeficiency virus infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human microtubulin 11 and encoded polynucleotide, applicable in diagnosis and treatment of e.g. developmental disorders, cancer, hemopathy, HIV infection, immunological diseases and various inflammations.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         human immunodeficiency virus infection; immunological disease; inflammation; embryonic development disorder; nervous system disorder; growth disorder; cytostatic; virucidal; immunodulatory; antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human;
human :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human microtubulin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         02-MAR-2000; 2000CN-00111820
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           immunological diseases, various inflammations, embryonic development disorders, disorders of the nervous system and growth disorders. The present sequence is an RT-PCR (reverse transcriptase PCR) primer used to isolate a nucleic acid encoding human microtubulin 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                                   Human plasminogen activator inhibitor 2-9.9 cDNA RT-PCR primer
                                                                                                                                                                                            27-FEB-2003
                                                                                                                                                                                                                                                                                                  ABS57470
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              645 CAGGCTGGAGTGCAGTGGCGCAAT 668
                                                                                                                                                                                                                                                                                                                                                                                                                                        24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BIOWINDOW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                     standard; DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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91.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              .2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1;
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Best Local S
Matches 22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mao
          Mao
                                                           06-NOV-2000;
                                                                                    06-NOV-2000; 2000CN-00127264.
                                                                                                             05-JUN-2002
                                                                                                                                     CN1352014-A.
                                                                                                                                                                Unidentified.
                                                                                                                                                                                     Starch precursor protein binding protein 13.42; Alzheimer's diseas tumour; development disorder; inflammation; immunological disease; haemopathy; HIV infection; cytostatic; anti-HIV; PCR; primer; ss.
                                                                                                                                                                                                                                                                   25-MAR-2003
                                                                                                                                                                                                                                                                                                                    ABZ21093 standard; DNA; 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 2; Page 17 (Disclosure); 33pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New human plasminogen activator inhibitor 2-9.9 polypeptide for treating e.g. hemorrhagic disease, thrombosis, cerebral infarction, various tumors, hemopathy, human immunodeficiency virus infection, and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                inflammations.
                                   (BODE-) BODE GENE DEV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (BODE-)
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                                                                                                                                                                                                                                         precursor
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                                                                                                                                                                                                                                                                                                                                                                                                          ACAGGCGTGAGCCACCACGCCCGG 896
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 2 A; 9 C; 9 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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                                                           2000CN-00127264.
                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                       protein binding
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                                   CO LTD SHANGHAI
                                                                                                                                                                                                                                                                                                                    BP.
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Pred. No. 1.
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                                                                                                                                                                                                                                         protein 13.42 PCR primer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1;
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                                                                                                                                                                                                                                          #2
                                                                                                                                                                                                               disease;
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WPI; 2003-381676/36

Reinhard C,

Walter A;

12-OCT-2001; 2001US-0328444P 11-OCT-2002; 2002WO-US032596

(CHIR) CHIRON CORP

17-APR-2003

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RESULT 423
ACA90126
                                                                                                                                                                                                                                                                                           W Human; ss; antisense; kinesin; CENP-E; Eg5; MCAK; colon cancer; stroke;

KW T cell cancer; B cell lymphoma; pancreatic cancer; breast cancer;

KW leukaemia; bladder cancer; stomach cancer; brain cancer; bone cancer;

KW oesophageal cancer; liver cancer; adrenalcarcinoma; lung cancer;

KW testicular cancer; heart cancer; ovarian cancer; uterine cancer;

KW head/neck cancer; cervical cancer; gall bladder cancer; spleen cancer;

KW parathrnoid cancer; penile cancer; gall bladder cancer; spleen cancer;

KW parathrnoid cancer; thyroid cancer; muscle cancer; stin cancer; melanoma;

KW myeloma sarcoma; teratocarcinoma; digestive cancer; sichaemia; epilepsy;

KW altoimmune disorder; viral infection; neurological disorder; meningitis;

KW liver disease; pancreatic disease; myocardial infarction; cerebral palsy;

KW liver disease; pancreatic disease; myocardial infarction; cerebral palsy;

KW amyotrophic lateral sclerosis; motor neuron disorder; multiple sclerosis;

KW creutzfeldt Jakob disease; muscular dystrophy; schizophrenia; amnesia;

KW Creutzfeldt mellitus; Grave's disease; cystic fibrosis; infection;

KW diabetic mellitus; Grave's disease; oystic fibrosis; infection;

KW myasthenia gravis; rheumatoid arthritis; osteoarthritis; scleroderma;

KW Sjogren's syndrome; systemic lupus erythematosus; toxic shock syndrome.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to starch precursor protein binding produced in the protein can be used for treating various disease, such as Alzheimer's disease, malignant tumours, development disorders, inflammations, immunological diseases, haemopathy and HIV infection. The present sequence is a PCR primer, which was used in an expectation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human kinesin gene(s) antisense oligonucleotide #9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ACA90126 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 24 BP; 3 A;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New starch precursor protein binding protein 13.42 polypeptide for treating e.g. Alzheimer's disease, malignant tumors, inflammations, immunological diseases, hemopathy and human immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2002-699445/76.
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Claim 5; Page 6; 57pp; English
                                                                                                                         Treatment of disease e.g. cancer, rheumatoid arthritis, Alzheimer's disease and Parkinson's disease involves administration of antisense
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cc administered either separately or in combination) and a pharmaceutical composition comprising the AS oligonucleotide and a carrier. The human ckinesin gene-targeting antisense oligonucleotides are useful for treatment of disease having aberrant cell proliferation such as cancer ce.g. colon cancer, T and B cell lymphoma, pancreatic cancer, breast concer, leukaemia, bladder cancer, stomach cancer, brain cancer, coesophageal cancer, liver cancer, adrenalcarcinoma, lung cancer, testicular cancer, heart cancer, ovarian cancer, brain cancer, head and neck cancer, thymus cancer, cervical cancer, gall bladder cancer, spleen cancer, thymus cancer, thyroid cancer, muscle cancer, gallal cancer, comdition associated with ischaemia and liver or pancreatic disease, cerebral neoplasm, Alzheimer's disease, extrapyramidal conder, myocardial infarction, stroke, epilepsy, ischaemic cerebrovascular conder, suppurative intracranial thrombophlebitis, multiple sclerosis, progressive neural muscular atrophy, retinitis pigmentosa, hereditary ataxia, suppurative intracranial thrombophlebitis, multiple sclerosis, subdural empyema, myelitis, paralysis, viral central nervous system contral retardation, cerebral palsv, autonomic nervous system disease, prion disease including kuru, Creutfeldt-Jakob disease, cerebral palsv, autonomic nervous system disease, cerebral palsv, autonomic nervous system disease, cerebral palsv, cerebral palsv, viral central nervous system contraction, cerebral palsv, viral central nervous system contraction, cerebral palsv, autonomic nervous system disease, paralysis, viral central nervous system disease, cerebral palsv, autonomic nervous system disease. mental retardation, cerebral paléy, autonomic nervous system disorder, muscular dystrophy, peripheral nervous system disorders, dermatomyositis, anxiety, schizophrenia, amesia, diabetic neuropathy, tratdive dyskinesia, Tourette's disease, cystic fibrosis, hypercholesterolaemia, diabetic mellitus, hyper- and hypoglycaemia, Grave's disease, neuralgia, Cushing's disease, Addison's disease, strointestinal disorders e.g. ulcerative colitis, duodenal ulcer, AIDS, allergic reactions, aucoimmune haemolytic anaemia, proliferative glomerulonephritis, inflammatory bowel disease, anaemia, proliferative glomerulonephritis, inflammatory bowel disease, Also included are the antisense oligonucleotides appearing as ACA90118-ACA90135, combination therapy involving administration of at least one chemotherapeutic or radionuclide and further involves administration of myasthenia gravis, rheumatoid arthritis, osteoarthritis, scleroderma, sjogren's syndrome, systemic lupus erythematosus, toxic shock syndrome, viral, bacterial, fungal, helminthic and protozoal infections. The present sequence is a human kinesin gene-targeting antisense oligonucleotide of the invention The invention relates to treatment of disease involving administering an antisense oligonucleotide. The oligonucleotide inhibits the expression of human kinesin gene. The human kinesin gene is CENP-E, human Eg5 or MCAK. at least one anti-sense oligonucleotide (the oligonucleotide is

Sequence 24 ₿P; 6 A; 11 C; 4 G; 3 T; 0 U; 0 Other;

Query Match Best Local Similarity Conservative 91.7%; 0; Mismatches Score 20.8; Pred. No. 1.2e+03 DB 1; 2 Length 24; , 0 Gaps

0;

밁 5 359 GCTCAAGCAGTCCACCTGCCTCAG 382 μ

SEXEXEXE RESULT 424 ACA90127 ACA90127 standard; DNA; 24 BP

10-JUL-2003 (first entry)

Human kinesin gene(s) antisense oligonucleotide #10

> KW Human; ss; antisense; kinesin; CENP-E; Eg5; MCAK; colon cancer; stroke;
> KW T cell cancer; B cell lymphoma; pancreatic cancer; breast cancer;
> KW leukaemia; bladder cancer; stomach cancer; brain cancer; bone cancer;
> KW desophageal cancer; liver cancer; adrenalcarcinoma; lung cancer;
> KW testicular cancer; heart cancer; ovarian cancer; uterine cancer;
> KW head/neck cancer; cervical cancer; ovarian cancer; uterine cancer;
> KW head/neck cancer; penile cancer; gall bladder cancer; spleen cancer;
> KW parathrnoid cancer; penile cancer; gall bladder cancer; skin cancer;
> KW thymus cancer; thyroid cancer; muscule cancer; skin cancer; melanoma;
> KW myeloma sarcoma; teratocarcinoma; digestive cancer; ischaemia; epilepsy;
> KW autoimmune disorder; viral infection; neurological disorder; meningitis;
> KW liver disease; pancreatic disease; mocardial infarction; cerebral palsy;
> KW Alzheimer's disease; Huntington's disease; parkinson's disease;
> KW amyotrophic lateral sclerosis; motor neuron disorder; multiple sclerosis;
> KW amyotrophic lateral sclerosis; motor neuron disorder; multiple sclerosis;
> KW creutzfeldt-Jakob disease; muscular dystrophy; schizophrenia; amnesia;
> KW diabetic mellitus; Grave's disease; cystic fibrosis; infection;
> KW diabetic mellitus; Grave's disease; pastrointestinal disorder;
> KW muschhonia cravis; rhanmatoid arthritis, orteoarthritis, sclerodarma. myasthenia gravis; rheumatoid arthritis; osteoarthritis; scleroderma; Sjogren's syndrome; systemic lupus erythematosus; toxic shock syndrome.

Homo sapiens

WO2003030832-A2

11-OCT-2002; 2002WO-US032596

12-OCT-2001; 2001US-0328444P

(CHIR) CHIRON CORP

Reinhard C, Walter A;

WPI; 2003-381676/36.

Treatment of disease e.g. cancer, rheumatoid arthritis, Alzheimer's disease and Parkinson's disease involves administration of antisense

Claim 5; Page 6; 57pp; English.

CC administered either separately or in combination and a pharmaceutical composition comprising the AS oligonucleotide and a carrier. The human C kinesin gene-targeting antisense oligonucleotides are useful for CC treatment of disease having aberrant cell proliferation such as cancer CC e.g. colon cancer, T and B cell lymphoma, pancreatic cancer, breast CC cancer, leukaemia, bladder cancer, stomach cancer, brain cancer, CC oesophageal cancer, liver cancer, stomach cancer, brain cancer, head and CC cancer, bone cancer, carrier cancer, uterine cancer, head and CC cancer, bone cancer, cancer, ovarian cancer, uterine cancer, head and CC cancer, thymus cancer, ponite cancer, grostate cancer, skin cancer, spleen CC parathrooid cancer, thyroid cancer, muscle cancer, sganglial cancer, CC lymphoma, autoimmune disorder, viral infection, neurological disorder, CC myocardial infarction, stroke, egilepsy, ischaemic cerebrovascular CC disease, cerebral neoplasm, Alzheimer's disease, perceptical disorder, amyotrophic lateral sclerosis, motor neuron disorders, corporatesive neural muscular atrophy. retinitis niomentosa. herefitary chemotherapeutic or radionuclide and further involves administration of at least one anti-sense oligonucleotide (the oligonucleotide is administered either separately or in combination) and a pharmaceutical human kinesin gene. The human kinesin gene is CENF-E, numan Ego or included are the antisense oligonucleotides appearing as ACA90118-ACA90135, combination therapy involving administration of at least one acceptance of the second of the seco The invention relates to treatment of disease involving administering an antisense oligonucleotide. The oligonucleotide inhibits the expression of human kinesin gene. The human kinesin gene is CENP-E, human Eg5 or MCAK. demyelinating disease, subdural empyema, myel: progressive neural muscular atrophy, retinitis pigmentosa, hereditary ataxia, suppurative intracranial thrombophlebitis, multiple sclerosis, demyelinating disease, bacterial and viral meningitis, brain abscess, subdural empyema, myelitis, paralysis, viral central nervous system myelitis, paralysis, viral central sease including kuru, Creutzfeldt-Ja head and

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RESULT 425
ACC57313
                                           Query Match
Best Local S
Matches 22
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Best Local :
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                                                                                                                                              Sequence
                                                                                                                                                                                                                Also disclosed are the polynuclectide encoding it, and a process for preparing the polypeptide using DNA recombination techniques. The application of the polypeptide is in treating diseases such as cancer human immunodeficiency virus (HIV) infection, The current sequence
                                                                                                                                                                                                                                                                                                                                                                 Example 2; Page 17 (disclosure); 33pp; Chinese.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Zinc finger protein; 11.55; human immunodeficiency virus; HIV; cancer; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Zinc finger protein 11.55 related PCR primer #SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        27-JUN-2003
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                                                                                                                                                                                         represents a zinc finger protein 11.55 related PCR
                                                                                                                                                                                                                                                                                                                The invention relates to a novel zinc finger protein designated 11.55
                                                                                                                                                                                                                                                                                                                                                                                                                 Polypeptide-zinc finger protein 11.55 and polynucleotide encoding
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198 CATGTTGGTCAGGCTGGTCTCGAA 221
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                                           l Similarity
22; Conser
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ilarity 91.7%;
Conservative
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                                                                                                                                         BP; 5 A;
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                                                                                                                                              6 C; 7 G; 6 T; 0 U; 0 Other;
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                                             0;
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                                           Score 20.8; D
Pred. No. 1.2e
0; Mismatches
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Pred. No. 1.
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                                                                                        Length 24;
                                             Indels
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ADG83872
                                                                 Query Match
Best Local
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                                                                                                                            The present invention describes a method for differentiating between ulcerative colitis and Crohn's disease based on the analysis of gene expression profiles in biopsy samples obtained from inflamed and optionally non-inflamed areas in the intestines of the patient. The method comprises determining the expression levels of at least two of a number of marker genes chosen from any of the 7 sequences SEQ ID NO:1 to 7 (see ADG83886, ADG83887, ADG83887, ADG83889, ADG83893) and ADG838921. The method can be used for differentiating between ulcerative colitis and Crohn's disease based on the analysis of gene expression profiles in biopsy samples obtained from inflamed and optionally non-inflamed areas in the intestines of the patient. The present sequence represents a PCR primer for a target genetic marker gene sequence which is used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                             Differentiating between ulcerative colitis and Crohn's disease based on the analysis of gene expression profiles in biopsy samples comprises determining the expression levels of at least two of a number of marker
                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                         Claim 7;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              04-SEP-2002;
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15-JUL-2002;
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18-JUL-2002;
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               862 GTGCTGGGATTACAGGCGTGAGCC 885
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                                                   l Similarity
22; Conserv
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2002SE-00002256.
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2002US-0407713P.
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2002SE-00002251.
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2002US-0395629P.
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                                                                                                       A; 4 C; 9 G; 6 T; 0 U; 0 Other;
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                                                                91.7%;
                                                                              2.1%;
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70
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                                                   Pred. No. 1.26); Mismatches
                                                                            Score 20.8;
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                                                                1.2e+03
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RESULT 428
ADB04739
ID ADB047
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AC ADB047
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AC ADB047
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DT 20-NOV
XX
DE Human
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Best Local Similarity
Matches 22; Conserv
                                                                                                                                                                                                                                                                        This oligonucleotide, termed primer CY6, was used with primer CY6B (see AAX24390) in a PCR amplification of novel human CC chemckine receptor CCR8 cDNA (see also AAX24385). Radiation hybrid mapping was performed by PCR using these primers. The CCR8 gene was mapped to human chromosome 3p22-p23. CCR8 (see AAW97868) is a G protein coupled receptor that plays an essential role in the membrane fusion step of HIV infection.

Establishment of stable, non-human cell lines and transgenic mammals having cells that coexpress CD4 and CCR8 provides valuable tools for research on HIV infection. Antibodies which bind to CCR8. CCR8 variants, and CCR8-binding agents capable of blocking membrane fusion between HIV and target cells represent potential anti-HIV therapeutics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                07-JUN-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    modulating immune responses or agents for the prevention or treatment HIV infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Chemokine receptor CCR8 PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAX24391 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              29-JUL-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                               Example 1; Page 53; 81pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Chemokine receptor; infection; therapy;
                                                                                      ADB04739 standard; DNA; 25
                                       20-NOV-2003
                                                                                                                                                                           864
               MDZ7
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                                                                                                                                                                           GCTGGGATTACAGGCGTGAGCCAC 887
                                                                                                                                                  GCTAGGATTACAGGCATGAGCCAC 24
             scanning oligonucleotide SEQ ID
                                                                                                                                                                                                                                                   BP; 8 A; 6 C; 7 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                    Conservative
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                                       (first entry)
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Pred. No. 1.2e+03;
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EP1281758-A2

sapiens

developmental

disorder; ss

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RESULT 429
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Best Local S
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                                                                                                                Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ7; MDZ12; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        vaccines. The present sequence was used to illustrate the
                                                                                                                                                                                                                    Human MDZ7 scanning
                                                                                                                                                                                                                                                                     20-NOV-2003
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ilarity 91.7%;
Conservative
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Pred. No. 1.2e
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Best Local Similarity
Matches 22; Conserv
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                                                                                                                                                                                                Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ7; MDZ12; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss.
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                                                                                                                                 EP1281758-A2
                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                Human MDZ7 scanning oligonucleotide SEQ ID 5724
                                                                                                                                                                                                                                                                                                                                                      ADB04738;
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                               02-AUG-2001; 2001US-00922181
                                                           30-JUL-2002; 2002EP-00016874
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(AEOM-) AEOMICA INC
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91.7%;
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Pred. No. 1.2e+03;
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Query Match
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                                                                                                                                 Sequence 25 BP; 8 A; 0 C; 4 G; 13 T; 0 U; 0 Other;
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                                                                                                                                                                                                             vaccines. The present sequence was used to illustrate the invention.
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2.1%;
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                          DB 1; Length 25;
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밁 S Matches Local 766 N Similarity ATTTTTTTGTATTTTTAGTAGAGA 789 **AATATTTTGTATTTTTAGTAGAGA** Conservative 0; Score 20.8; D Pred. No. 1.2e 0; Mismatches 25 1.2e+03; 2 Indels 0 Gaps

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ADB04740 RESULT 431 Human MDZ7 scanning oligonucleotide SEQ ID 5726. ADB04740 standard; DNA; (first entry)

developmental disorder; ss. Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ7; MDZ12; chromosome 7 chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer;

Homo sapiens.

EP1281758-A2

30-JUL-2002; 2002EP-00016874

02-AUG-2001; 2001US-00922181

(AEOM-) AEOMICA INC.

Gu Y, Nguyen

WPI; 2003-423107/40.

New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of of MDZ3

Example 8; SEQ MDZ4, MDZ7

ID NO 5726; 103pp;

English

or MDZ12, e.g. cancer

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RESULT 432
ADB04617
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Matches 22
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                                                                                                        New zinc finger-containing proteins and nucleic acide, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ7 or MDZ12, e.g. cancer.
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91.7%;
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The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ1, MDZ4, MDZ7, MDZ12. MDZ3 is encoded at chromosome 7g22.1, MDZ4 is encoded at chromosome fp21.3-22.2, MDZ7 is encoded at chromosome fp21.3-20.2).

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RESULT 433
ADB04578
ID ADB049
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The present invention relates to novel human zinc finger-containing proteins and their coding sequences; MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is encoded at chromosome 7q22.1, MDZ4 is encoded at chromosome 6p21.3-22.2, MDZ7 is encoded at chromosome fsp11.2 and MDZ12 is encoded at chromosome 1sq26.1. The MDZ3, MDZ4, MDZ7, and MDZ12 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12 The probes are
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                                                                                                                                                                                                                                                                                  New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ4, MDZ7 or MDZ12, e.g. cancer.
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Pred. No. 1.
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Best Local :
                                                                                proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is encoded at chromosome 7q22.1, MDZ4 is encoded at chromosome fp11.3-22.2, MDZ7 is encoded at chromosome fp11.2 and MDZ12 is encoded at chromosome 1sq26.1. The MDZ3, MDZ4, MDZ7, and MDZ12 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention
                                                     Sequence
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                                                                                                                                                                                                                                                                                                                                                                                            New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ4, MDZ7 or MDZ12, e.g. cancer.
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RESULT 435
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                                                                                                                                                                                                                                                   Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              invention relates to novel human zinc finger-containing
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                                                                                                                                                                                        DB 1;
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                                                                                                                                                                                     Length 25;
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RESULT 437
AAZ25152
ID AAZ251
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                                                                                                                                                                                    Query Match
Best Local
                                                                                                                                                                    Matches
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                                                                                                                                                                                                                                                                       more bases that are perfectly matching to a sequence anywhere of the first primer or the second primer, and the first primer at its 3' end does not contain eleven or more bases that are perfectly matching except one mismatch to a sequence anywhere of the first primer or the second primer. The method is useful for designing primers for simultaneous amplification of target DNA fragments in a single multiplex polymerase amplification of target DNA fragments in a single multiplex polymerase related to multifactorial diseases, the genome-scale detection of genetic alterations, the studies in pharmacogenetic reactions, the genotyping genetic polymorphisms in a large population, the gene expression profiling in various samples and high throughput genotyping technologies. This sequence corresponds to an example of a primer of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                amplification of target DNA fragments in a single multiplex polymerase chain reaction by aligning a first primer and a second primer. The method comprises: (a) aligning a first primer and a second primer; and (b) selecting the first primer where the first primer at its 3' end does not contain four or more bases that are perfectly matching to the 3' end sequence of the first primer or a second primer, the first primer at its 3' end does not contain seven or more bases that are perfectly matching except one mismatch to the 3' end sequence of the first primer or the second primer, the first primer at its 3' end does not contain six or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Designing primers for simultaneous amplification of target DNA fragments in a single multiplex polymerase chain reaction, for high throughput multiplex DNA sequence amplification, comprises aligning two primers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      07-OCT-2002; 2002US-0417009P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Single multiplex
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADO11741;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADO11741 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2004-340914/31.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Li H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           07-OCT-2003; 2003WO-US031874
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ss; primer; simultaneous amplification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 38; 120pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22-APR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO2004033649-A2
                                                                                                                                                                                                                                          Sequence 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to a method of designing primers for simultaneous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (UYNE-) UNIV NEW JERSEY MEDICINE & DENTISTRY
                                                                                                                               181
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                                                                                                                                                                l Similarity
22; Conserv
                                                                                                                               TAGAGATGGAGTTTCTCCATGTTG 204
                                                                                                                                                                                                                                          BP;
                                                                                                                                                                    Conservative
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                                                                                                                                                                                                                                          5 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PCR primer #1113.
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                                                                                                                                                                                  2.1%;
91.7%;
                                                                                                                                                                                                                                          4 C; 8 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        25
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                                                                                                                                                                  0; Mismatches
                                                                                                                                                                                                   Score 20.8;
                                                                                                                                                                                      Pred. No. 1
                                                                                              24
                                                                                                                                                                                      .2e+03
                                                                                                                                                                                                       DB 1;
                                                                                                                                                                                                   Length 25;
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                                                                                                                                                                    Gaps
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AAZ25152 standard; DNA; 22

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Human; short interspersed repetitive element; SINE; PCR; primer;

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RESULT 438
AAZ25149
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밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention describes a restriction primer for eukaryotic short conterspersed repeated sequences (SINE), which has one or more additional CC bases that are a mismatch to, or are unrelated to, the 3'-terminal end of CC the SINE. The annealing temperature of the primer to the DNA sequence is CC chain reaction (PCR). The PCR fragments obtained are subjected to CC cletrophoresis to obtain a fingerprint. By comparing the polymerase CC distinguished. The primer is used for amplifying a eukaryotic deoxyribonucleic acid (DNA) sequence, pinched between SINE sequences by CC clustions and ecological studies. DNA specimens in traces conspicultural identification of humans for medical and legal CC applications and ecological studies. DNA specimens in traces CC februarity in mass) can be used for individual discrimination CC (approximately 10 ng in mass) can be used for individual discrimination constitution of humans for medical and legal constitutions and ecological studies. DNA specimens in traces constitution of humans for medical and legal constitution of humans for medical and legal
                                                                                                                                                                                                                                                                                                                                                                                                 Matches
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Homo sapie
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Restriction primer for distinguishing individuals with short interspersed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     eukaryote; restricted polymerase chain reaction fidentification; DNA specimen; discrimination; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAZ25152;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 22 BP; 6 A; 5 C; 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 6; Page 3; 17pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1999-583348/50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10-JUL-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; short interspersed repetitive element; SINE; PCR; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        13-DEC-1999
                                       Human short
                                                                                         13-DEC-1999
                                                                                                                                                                                    AAZ25149 standard; DNA; 22 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             of eukaryotes using the primer in a polymerase chain reaction
AAZ25143 to AAZ25191 represent specifically claimed examples of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 repeated sequence of eukaryotes by restricted polymerase chain reaction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (NORQ ) NORINSUISANSHO SUISANCHO YOSHOKU KENKYUSHOCHO
                                                                                                                                        AAZ25149
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                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            present invention
                                                                                                                                                                                                                                                                                                        GGATTACAGGCGTGAGCCACTA 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18; restriction primer; short interspersed repeated sequence;
restricted polymerase chain reaction fingerprinting;
                                          interspersed repetitive element PCR primer #7.
                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                         (first entry
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                                                                                                                                                                                                                                                                                                                                                                                                                        2.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 20.4;
                                                                                                                                                                                                                                                                                                                                                                                                                             Pred.
                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                             No.
                                                                                                                                                                                                                                                                                                                                                                                                                             1.1e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1; Length 22;
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Best Local
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ANZESI43 to ANZESI41 represent specifically claimed examples of primers
JP2913035-B1
                                        Homo
                                                                                                                    Human; short interspersed repetitive element; SINE; PCR; primer; Oncorhynchus; restriction primer; short interspersed repeated sequence; eukaryote; restricted polymerase chain reaction fingerprinting;
                                                                                                                                                                                                      Human short interspersed repetitive element PCR primer #4.
                                                                                                                                                                                                                                                 13-DEC-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention describes a restriction primer for eukaryotic short interspersed repeated sequences (SINE), which has one or more additional.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim
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                                                                                                                                                                                                                                                                                                                                 AAZ25146 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (NORQ ) NORINSUISANSHO SUISANCHO YOSHOKU KENKYUSHOCHO
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                                                                                                     identification;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
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                                                                                                                                                                                                                                             (first entry)
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                                                                                                     DNA
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                                                                                                                                                                                                                                                                                                                                 DNA;
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95.5%;
                                                                                                                                                                                                                                                                                                                                   22
                                                                                                                                                                                                                                                                                                                                   ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 20.4;
Pred. No. 1.
                                                                                                     discrimination;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1.1e+03;
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AAC69376/c
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Best Local S
Matches 21
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                   15-MAR-2000; 2000WO-IB000532
                                                            21-SEP-2000.
                                                                                                     WO200055318-A2
                                                                                                                                            Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                      Human ABC1
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CC bases that are a mismatch to, or are unrelated to, the 3'-terminal end of the SINE. The annealing temperature of the primer to the DNA sequence is kept higher than the fusion temperature of the primer during polymerase chain reaction (PCR). The PCR fragments obtained are subjected to electrophoresis to obtain a fingerprint. By comparing the polymorphs from the electrophoresis band pattern, eukaryotic individuals are distinguished. The primer is used for amplifying a eukaryotic deoxyribonucleic acid (DNA) sequence, pinched between SINE sequences by colymerase chain reaction (PCR) fingerprinting. In particular it may be used individual identification of humans for medical and legal applications and ecological studies. DNA specimens in traces (approximately 10 mg in mass) can be used for individual discrimination of eukaryotes using the primer in a polymerase chain reaction (PCR).

ARZ25143 to ARZ5191 represent specifically claimed examples of primers
Human ABC1 cholesterol transporter; currently density transporter; high density transporter; high density transporter disease; TD; familial HDL deficiency; FHA; polymorphism; cardiovascular disease; coronary artery disease; coronary restent cerebrovascular disease; peripheral vascular disease; cerebrovascular disease; Niemann-Pick disease; Huntington's disease; Alzheimer's disease; Niemann-Pick disease; gene therapy; genetic dia
                                                                                                                                                                                                                                Human ABC1 cholesterol transporter; chromosome 9q31;
ATP-binding cassette; HDL deficiency disorder; high density lipoprotein;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Restriction primer for distinguishing individuals with short interspersed repeated sequence of eukaryotes by restricted polymerase chain reaction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (NORQ ) NORINSUISANSHO SUISANCHO YOSHOKU KENKYUSHOCHO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention describes a restriction interspersed repeated sequences (SINE), which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              868 GGATTACAGGCGTGAGCCACCA 889
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GGATTACAGGCGTGAGCCACAA 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22 BP; 7 A; 5 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      present invention
                                                                                                                                                                                                                                                                                                                                              BAC contig polymorphic site, SEQ ID NO:275
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         98JP-00195692
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 20.4;
Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1.1e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         primer for eukaryotic short
has one or more additional
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                               restenosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
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CC disease, particularly coronary artery disease, but also cerebrovascular CC disease, coronary restenosis, and peripheral vascular disease. CC conversely, a high level of HDI has protective effects against CC cardiovascular disease. The invention provides genetic constructs and CC transgenic cells and non-human animals comprising human ABC1 nucleic cardiovascular disease comprising the administration of a revention of cc cardiovascular disease comprising the administration of an expression CC vector encoding ABC1 or an active fragment thereof. The invention also encompasses compounds which mimic ABC1 activity, compounds which CC stimulate ABC1 expression and methods of screening for such compounds. It further relates to methods for determining whether a patient has an CC increased risk for cardiovascular disease due to polymorphisms in the ABC1 gene. Human ABC1 proteins and nucleotides can be used to treat or prevent cardiovascular disease, especially coronary artery disease, cerebrovascular disease, especially coronary artery disease, disease. They may also be used in the treatment of diseases associated disease, Huntington's disease, X-linked adrenoleukodystrophy and cancer. The invention specifically excludes proteins with the exact amino acid sequences of Gendak Accession No: CAN10005.1 and X75926, and the nucleic acid with the exact remnerce as Cannark Accession No: An010376 1 The
RESULT 441
AAF74132
ID AAF741
XX
AC AAF741
XX
AC AAF741
XX
DT 30-APR
XX
DT 30-APR
XX
DE Primer
                                                                                                                                                                                                                                      В
                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local
                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 15-MAR-1999;
08-JUN-1999;
17-JUN-1999;
01-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to the human ABC1 cholesterol transporter protein (B38082) and to nucleic acid sequences (C69120) which encode it. ABC1 is a member of the ATP-binding cassette (ABC transporter) superfamily of proteins, and plays a crucial role in cholesterol transport, particularly intracellular cholesterol trafficking in monocytes and fibroblasts, being involved in cholesterol trafficking in monocytes and fibroblasts, being involved in cholesterol efflux from the cell. The gene encoding ABC1 is located on chromosome 9931, and mutations in this gene are associated with two genetic HDL (high density lipoprotein) deficiency disorders, Tangier disease (TD) and familial HDL deficiency (FHA). These diseases are distinguishable in that TD is an autosomal recessive disorder, while
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hayden
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FHA is inherited as an autosomal dominant trait. Low levels of HDL ("good cholesterol") in the blood correlate with a high risk of cardiovascular disease, particularly coronary artery disease, but also cerebrovascular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New ABC1 polypeptide is us
biological activity, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2000-587528/55
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example; Fig 11; 229pp; English
                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (UYBR-)
                                            30-APR-2001
                                                                                       AAF74132;
                                                                                                                               AAF74132 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      d with the exact sequence as GenBank Accession No: AJ012376.1. The sent sequence represents a polymorphic site of the human ABC1 gene
                                                                                                                                                                                                                                                                              533
                                                                                                                                                                                                                                    22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         VIND
                                                                                                                                                                                                                                                                                                                         l Similarity
21; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                    sequence
                                                                                                                                                                                                                                                            TCCTCCTGCCTCAGCCTCCCAA 554
                                                                                                                                                                                                                                        TTCTCCTGCCTCAGCCTCCCAA 1
                                                                                                                                                                                                                                                                                                                                                                                                            BP; 6 A;
                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BRITISH COLUMBIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   99US-0124702P.
99US-0138048P.
99US-0139600P.
99US-0151977P.
                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                    represents a polymorphic
                                                                                                                                 DNA;
                                                                                                                                                                                                                                                                                                                                           2.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   is useful for treating diseases associated with ABC1 e.g. Alzheimer's disease, Huntington's disease and
                                                                                                                                                                                                                                                                                                                                                                                                            2 C; 11 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pimstone
                                                                                                                                 22
                                                                                                                                 ВP
                                                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                                                               Pred.
                                                                                                                                                                                                                                                                                                                                                                   Score 20.4;
                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SN;
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                                                                                                                                                                                                                                                                                                                                                 .1e+03
                                                                                                                                                                                                                                                                                                                                                                   DB 1; Length 22;
                                                                                                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                                                                                         Gaps
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Matches
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New isolated polynucleotide comprising a polymorphic variant for the solute carrier family 6 neurotransmitter transporter, serotonin member 4 gene for identifying drugs for treating disorders related to expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Solute carrier fami
genotyping; allele
                                                                                                                                                                                                                                                                                                                                                                                                             genotyping oligonucleotide is used to detect a polymorphism in the SLC6A4
                                                                                                                                                                                                                                                                                                                                                                                                                                        serotonin member 4 (SLC6A4) gene or a fragment of it or a sequence complementary to the first sequence. The invention is used in produce recombinant organism that can be used to express SLC6A4 for protein structure analysis and binding studies. A composition comprising a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             sequence for the solute carrier family 6 neurotransmitter transporter, serotonin member 4 (SLC6A4) gene or a fragment of it or a sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       31-JUL-2000; 2000WO-US020638
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     08-FEB-2001.
                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to a polymorphic variant of a reference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 1; Page 38; 152pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2001-123317/13.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
                                             ADL66997 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                            gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Denton RR,
ADL66997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (GENA-) GENAISSANCE PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   the protein.
                                                                                                                                                                                                           870 ATTACAGGCGTGAGCCACCACG 891
                                                                                                                                                                                                                                                         21;
                                                                                                                                                              1 ATTACAGGTGTGAGCCACCACG 22
                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Duda A,
                                                                                                                                                                                                                                                                                                                                            BP; 6 A; 6 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           family 6 neurotransmiter transporter;
lele specific oligonucleotide; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            99US-0146290P.
                                                                                                                                                                                                                                                                              2.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nandabalan
                                                  ₽P
                                                                                                                                                                                                                                                           0; Mismatches
                                                                                                                                                                                                                                                                                   Pred.
                                                                                                                                                                                                                                                                                                      Score 20.4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ۲
                                                                                                                                                                                                                                                                                   No. 1.1e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sanchis A,
                                                                                                                                                                                                                                                                                                      DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Stephens
                                                                                                                                                                                                                                                                                                    Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        sectonin 4;
                                                                                                                                                                                                                                                           <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          producing
                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SLC6A4;
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RESULT 442
ADL66997
05-SEP-2002;
19-NOV-2002;
                                              05-SEP-2003; 2003WO-US027705
                                                               18-MAR-2004
                                                                                                             DNA polymerase; anti-DNAP antibody; reverse transcriptase; anti-RT antibody; single strand binding protein; SSB; ss;
                                                                                                                                      Multiplex PCR primer
                                                                                                                                                      03-JUN-2004
         (INVI-) INVITROGEN CORP
                                                                               WO2004022770-A2
                       2002US-0408609P
2002US-0427867P
                                                                                                                                                      (first entry)
                                                                                                                                       #1.
                                                                                                               primer.
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we colon disorder; intestinal disorder; cardiovascular disorder;

we muscular disorder; blood disorder; immune disorder; bone disorder;

we joint disorder; metabolic disorder; nutritive disorder; cancer;

we kidney disorder; liver disorder; puscate disorder; breast disorder;

we vary disorder; stomach disorder; pancreas disorder; spleen disorder;

we kin disorder; stomach disorder; pancreas disorder; spleen disorder;

we kin disorder; thyroid disorder; antiparkinsonian; antimanic;

we thymus disorder; thyroid disorder; antiparkinsonian; antimaric;

we cytostatic; antiinflammatory; vasotropic; antidiarrhecic; antidiabetic;

we CNS; central nervous system; respiratory; antidiarrhecic; antidiabetic;

we contral nervous system; respiratory; antidiarrhecic;

we contr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a new composition which comprises at least one anti-DNA polymerases (anti-DNAP) antibody and/or at least one anti-reverse transcriptase (anti-RT) antibody, and at least one single strand binding protein (SSB) or at least two different SSBs. The compositions are useful for nucleic acid synthesis reactions or are generated during nucleic acid synthesis reactions or are generated during one or more nucleic acid molecules. The compositions and methods are also be used in amplifying nucleic acid molecules, in reverse transcription/amplification. The present sequence is used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 22 BP; 4 A; 3 C; 10 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 4; Page 89; 201pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New compositions comprising one or more anti-reverse transcriptase antibodies, anti-DNA polymerases or single strand binding proteins, useful for synthesizing nucleic acids.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2004-248479/23
Gaitanaris GA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human novel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADO30457 standard; DNA; 22
                                                                                                          09-SEP-2002; 2002US-0409303P
09-APR-2003; 2003US-0461329P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    29-JUL-2004
                                                        (PRIM-) PRIMAL
                                                                                                                                                                                         09-SEP-2003; 2003WO-US028226
                                                                                                                                                                                                                                                                                                   WO2004040000-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       protein-coupled receptor; GPCR; drug screening; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     647 GGCTGGAGTGCAGTGGCGCAAT 668
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GGCTGGAGTGCAGTGGTAAT 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GPCR PGR4 RT-PCR primer, SEQ ID NO:1560
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                        INC.
Bergmann JE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2.1%;
95.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 20.4;
Pred. No. 1
Gragerov A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1.1e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1;
  Hohmann J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 22
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  শ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
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Polymorphism; allergic disea

disease;

PCR primer; ss.

human;

interleukin 4 receptor-alpha; IL4R-alpha;

Homo sapiens WO200104270-A1

Human IL4Ralpha gene PCR primer #84.

18-APR-2001

(first entry)

AAF69748 standard; DNA; 23

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GTTGGTCAGGTTGGTCTCGAAC 22

Matches Query Match

21;

Conservative

0; Mismatches Score 20.4; Pred. No. 1.

1.1e+03 DB 1;

Length 22; Indels

0,

Gaps

0

Local Similarity

2.1%; 95.5%;

201 GTTGGTCAGGCTGGTCTCGAAC 222

Sequence

22 BP; 3 A; 4 C; 8 G; 7 T; 0 U; 0 Other;

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CC diseases including neurological disorders (e.g., Albheimer's disease, CC depression, diabetic neuropathy, Parkinson's disease or schizophrenia); CC disorders of the adrenal gland; disorders of the colon or intestine CC (e.g., Crohn's disease, diarrhoea, food poisoning or irritable bowel CC syndrome); cardiovascular disorders (e.g., angina, cardiac arrhythmia or CC myocardial infarction); muscular disorders, blood disorders (e.g., anaemia or leukaemia); immune disorders (e.g., autoimmune disorders or CC Albb); bone and joint disorders (e.g., osteoarthritis, rheumatoid CC arthritis, gout or osteoporosis); metabolic or nutritive disorders (e.g., obesity, enzyme deficiency-related diseases or vitamin deficiency-related CC diseases); and disorders of the kidney, liver, lung, breast, ovary, CC uterus, prostate, testis, skin, stomach, pancreas, spleen, thymus and CC used in the isolation of cDNA encoding the novel human GPCR PGR4. Note: CT thyroid (e.g., cancers). The present sequence represents a PCR primer CC used in the isolation of cDNA encoding the novel human GPCR PGR4. Note: CT entil sequence data for this pattent did not form part of the printed CC specification; those sequences not shown were obtained in electronic CC format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (GPCRS) and nucleic acids encoding them. The invention also relates to sequences at least 90% identical to the GPCR proteins and nucleic acids of the invention, methods of tracting, preventing or diagnosing diseases associated with GPCRs of the invention; methods of screening for compounds useful in the treatment of GPCR-related diseases; a transgenic mouse comprising a GPCR gene of the invention; a mouse comprising a mutation in a GPCR transgene or in an endogenous GPCR gene; calls derived from the trasngenic mice; kits comprising several mice, each of which has a mutation in a different GPCR gene of the invention, and kits comprising probes which hybridise to GPCR polymucleotides of the invention. The invention further discloses variants of the GPCR polypeptides and vectors comprising a GPCR nucleic acid. The GPCR nucleic acids and proteins may be used in the diagnosis, treatment or prevention of a wide variety of discase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to human and mouse G protein-coupled receptors (GPCRs) and nucleic acids encoding them. The invention also relates
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel mammalian G protein coupled receptors, useful for identifying compounds that modulates diagnosing and treating disease condition associated with GPCR dysfunction e.g. autoimmune diseases, angina
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           pectoris, Parkinson's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Madisen L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID NO 1560; 542pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mcilwain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ŗ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pavlova MN,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Vassilatis D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Zeng
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RESULT 445
AAH49787
ID AAH497
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Best Local S
Matches 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention relates to polymorphisms of the human interleukin 4 receptor-alpha gene (IL4R-alpha; see AAF57718 for the reference sequence). Polymucleotides comprising polymorphic gene variants are useful for therapeutic purposes. For example, where a patient may benefit from expression of a particular IL4Ralpha protein isoform, an expression vector encoding the isoform may be administered to the patient. It may desirable to decrease or block expression of a particular IL4Ralpha isogene, which may be done by turning off by transforming a targeted organ, tissue or cell population with an expression vector that expresses high levels of untranslatable mRNA for the isogene. Specific therapeutics identified by these methods may be useful for allergic diseases. The present sequence is a PCR primer for human IL4R-alpha
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              13-JUL-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New isolated polynucleotide useful for the identification in allergic diseases is new.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2001-103078/11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Windemuth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Chew
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 1;
           Uncoiling enzyme 9 and encoded polynucleotide, applicable in diagnosis and treatment of malignant tumor, hemopathy, HIV infection, immunologidiseases and various inflammation.
                                                                                                                                     24-DEC-1999;
                                                                                                                                                             18-DEC-2000; 2000WO-CN000616
                                                                                                                                                                                                               WO200149860-A1
                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                              immunological
                                                                                                                                                                                                                                                                            Human;
                                                                                                                                                                                                                                                                                                      Human uncoiling enzyme 9 coding sequence PCR primer
                                                                                                                                                                                                                                                                                                                               25-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                  AAH49787
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (GENA-)
                                                             WPI; 2001-432884/46.
                                                                                                             (BIOW-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       578
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                                                                                                                                                                                                                                                                uncoiling enzyme
logical disease; i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
21; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENAISSANCE PHARM INC
                                                                                     Xie
                                                                                                             BIOWINDOW GENE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Denton RR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CCACTACACCTGGCTAATTTTT 599
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    23
                                                                                                                                                                                                                                                                                                                                                                                  standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                CCACCACACCTGGCTAATTTTT 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Page 64; 188pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2000WO-US019094.
                                                                                                                                                                                                                                                                                                                               (first entry)
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                                                                                                                                      99CN-00125756
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2.1%;
95.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    8 C; 3 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                inflammation;
                                                                                                              DEV LTD
                                                                                                                                                                                                                                                                               9
                                                                                                                                                                                                                                                                                                                                                                                  24
                                                                                                                                                                                                                                                                              cancer; haemopathy; HIV infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 20.4;
Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nandabalan
                                                                                                              SHANGHAI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1.2e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 1;
                                                                                                                                                                                                                                                                therapy;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     of therapeutics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
                      immunological
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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TGGAGTCTCACTCTGTTACCCA

925

TGGAATCTCACTCTGTTACCCA 946

Query Match Best Local S

21;

Conservative

<u>,</u>

Mismatches

Similarity

2.1%; 95.5%;

Pred. No. 1.2e+03

DB 1;

Length

Indels

<u>;</u>

Gaps

0,

Sequence 24

BP; 4 A;

7 C; 6 G; 7 T; 0 U; 0 Other;

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RESULT 446
ABS56869
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Query Match
Best Local S
Matches 21
                      The invention relates to the human receptor related tyrosine kinase 10.01, a polymucleotide encoding the polypeptide and a method for producing the polypeptide by DNA recombination technology. The polypeptide is used for curing several diseases such as embryonic development deformity, tumours, diabetes, menstrual disorder, peptic ulcer, anaemia and epilepsy. This sequence represents a reverse transcriptase PCR (RT-PCR) primer used in isolation of cDNA encoding human receptor related tyrosine kinase 10.01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention provides the protein and coding sequences of human uncolling enzyme 9. The sequences can be used in the treatment of cancer, haemopathy, HIV infection, immunological diseases and inflammation. The present sequence is a PCR primer for the coding sequence of the invention
                                                                                                                                                                                         New polypeptide-human receptor related tyrosine kinase 10.01 for treating embryonic development deformity, tumor, diabetes, menstrual disorder, peptic ulcer, anemia and epilepsy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cancer; anaemia; epilepsy; RT-PCR; reverse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; receptor related tyrosine kinase 10.01; primer; ss; peptic ulcer; embryonic development deformity; tumour; diabetes; menstrual disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human receptor related tyrosine kinase 10.01 cDNA RT-PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 24 BP; 5 A; 4 C; 7 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 3; Page 11; 32pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                  29-SEP-2000; 2000CN-00125572
                                                                                                                                                                                                                                                                                                                                                                                                         24-APR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                        CN1345961-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30-JAN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABS56869
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABS56869 standard;
                                                                                                                                                                                                                                                       WPI; 2002-539360/58.
                                                                                                                                                                                                                                                                                    Mao Y,
                                                                                                                                                                                                                                                                                                                                                                             29-SEP-2000; 2000CN-00125572
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                             Example 2; Page 18 (Disclosure); 34pp; Chinese
                                                                                                                                                                                                                                                                                                                  (SHAN-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  182 AGAGATGGAGTTTCTCCATGTT 203
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ب
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                                                                                                                                                                                                                                                                                    Xie Y;
                                                                                                                                                                                                                                                                                                                  SHANGHAI BIOWINDOW GENE DEV INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AGAGATGGAGTTTCGCCATGTT 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          95.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              <u>,</u>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20.4;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
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22 TCCTGCCTCAGCCTCCCCAGTA 1

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RESULT 447
AAH40563/c
                                                                                                                                                                                                          SNP flanking sequence, the SNPE primer is used as a genotyping primer.

The oligonucleotides are useful for genotyping a nucleic acid sample by performing a single-nucleotide primer extension reaction. The oligonucleotides are useful for determining the presence, absence or claiming to a secondary and for genotyping nucleic acid samples, for e.g. to assess by association analysis the genotype of an individual or group of individuals, having a pathological phenotypic trait suspected of being caused by one or more SNPs. Phenotypic traits include diseases e.g. agammaglobulinaemia, diabetes insipidus, Lesch-Nyhan syndrome, muscular dystrophy, familial hypercholesterolaemia, polycystic kidney disease, osteogenesis imperfecta and acute intermittent porphyria. Phenotypic traits also include symptoms of or susceptibility to multifactorial disease of which a component is or may be generic such as autoimmune disease, including, rheumatoid arthritis, multiple sclerosis, inflammation, cancer, nervous system diseases and infection by pathogenic microorganism. The method is also useful in forensic investigations and paternity analysis. The present sequence represents a single mucleotide primer extension (SNPE) primer specific for a human SNP containing DNA
                                                                    Matches
                                                                                     Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide primer extension (SNPB) primers, and the sequences of regions flanking sites of single nucleotide polymorphisms SNPs. The present invention includes kits for determining the presence or absence of a SNP, using the oligonucleotides of the invention. The PCR primers are used to amplify a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nuc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SNP specific SNPE primer SEQ ID
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                                                                                                                                                   Sequence 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1; Page 67; 83pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Picoult-Newburg L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15-OCT-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (ORCH-) ORCHID
536 TCCTGCCTCAGCCTCCCAAGTA 557
                                                                                          Similarity
                                                                                                                                                     BP; 7
                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    forensic investigation; paternity analysis; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BIOSCIENCES INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       99US-0160096P.
                                                                                                                                                   A; 4 C; 11 G; 3 T; 0 U; 0 Other;
                                                                                 2.1%;
95.5%;
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                                                                 0;
                                                                                     Score 20.4;
Pred. No. 1.
                                                                 Mismatches
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                                                                                     .2e+03
                                                                                                       DB 1; Length 25;
                                                                    Indels
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                                                                 Gaps
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RESULT 448
AAH38991/c
ID AAH389
XX
   FARRARA AND COLORO COLO
                                                                             CC primer extension (SNPE) primers, and the sequences of regions flanking CC sites of single nucleotide polymorphisms SNPs. The present invention CC includes kits for determining the presence or absence of a SNP, using the coligonucleotides of the invention. The PCR primers are used to amplify a CC SNP flanking sequence, the SNPE primer is used as a genotyping primer. CC The oligonucleotides are useful for genotyping a nucleic acid sample by performing a single-nucleotide primer extension reaction. The CC oligonucleotides are useful for determining the presence, absence or CC identity of a SNP and for genotyping nucleic acid sample by cases by association analysis the genotypic trait suspected of being CC caused by one or more SNPs. Phenotypic traits include diseases e.g. CC agammaglobulinaemia, diabetes insipidus, Lesch-Nyhan syndrome, muscular CC diseases imperfecta and acute intermittent porphyria. Phenotypic traits also include symptoms of or susceptibility to multifactorial CC diseases, including, rheumatoid arthritis, multiple sclerosis, clinflammation, cancer, nervous system diseases and infection by pathogenic microorganism. The method is also useful in forensic investigations and cc semmence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequences AAH37205 - AAH40944 represent PCR primers,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 1; Page 59; 83pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               acid sample.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nucleic
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Sequence
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      4 T; 0 U;
      0 Other;
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Query Match Best Local S Matches 22

Local Similarity

Conservative

88.0%;

2.0%;

Score 20.2; Pred. No. 1.

1.3e+03;

Length 25; Indels

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Gaps

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CCCGCCTTGACCTCCCAAAGTGCTG 1 CCTGCCTCGGCCTCCCAAAGTGCTG

867

843 25

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RESULT 449
AAH40899
CC SNP flanking sequence, the SNPE primer is used as a genotyping primer.

CC The oligonucleotides are useful for genotyping a nucleic acid sample by reforming a single-nucleotide primer extension reaction. The coligonucleotides are useful for determining the presence, absence or identity of a SNP and for genotyping nucleic acid samples, for e.g. to casses by association analysis the genotype of an individual or group of individuals, having a pathological phenotypic trait suspected of being caused by one or more SNPs. Phenotypic traits include diseases e.g. cagammaglobulinaemia, diabetes insipidus, Lesch-Nyhan syndrome, muscular dystrophy, familial hypercholesterolaemia, polycystic kidney disease, consteogenesis imperfecta and acute intermittent porphyria. Phenotypic traits also include symptoms of or susceptibility to multifactorial confideases, including, rheumatoid arthritis, multiple sclerosis, including, rheumatoid arthritis, multiple sclerosis, cancer, nervous system diseases and infection by pathogenic confidentity analysis. The present sequence represents a single nucleotide primer extension (SNPE) primer specific for a human SNP containing DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  primer extension (SNPE) primers, and the sequences of regions flanking sites of single nucleotide polymorphisms SNPs. The present invention includes kits for determining the presence or absence of a SNP, using the oligonucleotides of the invention. The PCR primers are used to amplify a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15-OCT-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1; Page 68; 83pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide
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Sequence

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BP; 3

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G; 7 T; 0 U; 0 Other; Score 20.2; DB 1; Pred. No. 1.3e+03;

Length 25

Sequence 25 BP;

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Local Similarity

2.0**%;** 88.0**%**; 9 C; 6

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RESULT 450
AAH37979/c
ID AAH379
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cc includes kits for determining the presence of a SNP, using the coligonucleotides of the invention. The PCR primers are used to amplify a cc SNP flanking sequence, the SNPE primer is used as a genotyping primer. CC The oligonucleotides are useful for genotyping a nucleic acid sample by coligonucleotides are useful for determining the presence, absence or coligonucleotides are useful for determining the presence, absence or coligonucleotides are useful for determining the presence, absence or coligonucleotides are useful for determining the presence, absence or coligonucleotides are useful for determining the presence, absence or coligonucleotides are useful for determining the presence, absence or coligonucleotides are useful for determining the presence, absence or coligonucleotides an analysis the genotype of an individual or group of individuals, having a pathological phenotypic traits usspected of being caused by one or more SNPs. Phenotypic traits include diseases e.g. cagammaglobulinaemia, diabetes insipidus, Lesch-Nyhan syndrome, muscular cystrophy, familial hypercholesterolaemia, polycystic kidney disease, costeogenesis imperfecta and acute intermittent porphyria. Phenotypic craits also include symptoms of or susceptibility to multifactorial contisease of which a component is or may be genetic such as autoimmune diseases, including, rheumatoid arthritis, multiple sclerosis, contenned to the present sequence represents a single nucleotide primer extension (SNPE) primer specific for a human SNP containing DNA containing DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    primer extension (SNPE) primers, and the sequences of regions flanking sites of single nucleotide polymorphisms SNPs. The present invention includes kits for determining the account.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis;
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                            Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide CC primer extension (SNPB) primers, and the sequences of regions flanking CC sites of single nucleotide polymorphisms SNPs. The present invention CC includes kits for determining the presence or absence of a SNP, using the CC SNP flanking sequence, the SNPE primer is used as a genotyping primer. CC The oligonucleotides are useful for genotyping a nucleic acid sample by CC performing a single-nucleotide primer extension reaction. The CC performing a single-nucleotide primer extension reaction. The CC cligonucleotides are useful for determining the presence, absence or CC identity of a SNP and for genotyping nucleic acid samples, for e.g. to CC assess by association analysis the genotype of an individual or group of CC cligonucleotides are useful for determining the presence, absence or CC dystrophy, familial hypercholesterolaemia, polycystic kidney disease, c.g. caused by one or more SNPs. Phenotypic trait suspected of being CC agammaglobulinaemia, diabetes insipidus, Leech-Nyhan syndrome, muscular CC dystrophy, familial hypercholesterolaemia, polycystic kidney disease, c.g. categenesis imperfecta and acute intermittent porphyria. Phenotypic traits also include symptoms of or susceptibility to multifactorial CC diseases, include symptoms of or susceptibility to multifactorial constants. The method is also useful in forensic investigations and cC paternity analysis. The present sequence represents a single nucleotide primer extension (SNPE) primer specific for a human SNP containing DNA
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Pred. No. 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 452
AAH39587
                          performing a single-nucleotide primer extension reaction. The composition of the personal production and for genotyping nucleic acid samples, for e.g. to identity of a SNP and for genotyping nucleic acid samples, for e.g. to casess by association analysis the genotype of an individual or group of individuals, having a pathological phenotypic trait suspected of being caused by one or more SNPs. Phenotypic traits include diseases e.g. cagammaglobulinaemia, diabetes insipidus, Lesch-Nyhan syndrome, muscular dystrophy, familial hypercholesterolaemia, polycystic kidney disease, osteogenesis imperfecta and acute intermittent porphyria. Phenotypic traits also include symptoms of or susceptibility to multifactorial disease of which a component is or may be genetic such as autoimmune inflammation, cancer, nervous system diseases and infection by pathogenic microorganism. The method is also useful in forensic investigations and
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis;
                                                                                                                                                                                                                                                                                                                            primer extension (SNPE) primers, and the sequences of regions flanking sites of single nucleotide polymorphisms SNPs. The present invention includes kits for determining the presence or absence of a SNP, using the oligonucleotides of the invention. The PCR primers are used to amplify a
                                                                                                                                                                                                                                                                                         SNP flanking sequence, the SNPE primer is used as a genotyping primer. The oligonucleotides are useful for genotyping a nucleic acid sample by
                                                                                                                                                                                                                                                                                                                                                                                                        Sequences AAH37205
                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1; Page 62; 83pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nuc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2001-290930/30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SNPE primer SEQ ID 2383
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   forensic investigation; paternity analysis; primer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA;
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Pred. No. 1.3e+03
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hy; familial hypercholesterolaemia;
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primer extension (SNPE)

paternity analysis. The present sequence represents

primer specific

for

a single nucleotide

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RESULT 453
AAH39123
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Crimer extension (SNPE) primers, and the sequences of regions flanking concludes kits for determining the presence or absence of a SNP, using the coligonucleotides of the invention. The PCR primers are used to amplify a CC SNP flanking sequence, the SNPE primer is used as a genotyping primer. CC performing a single-nucleotide portiner extension reaction. The CC performing a single-nucleotide primer extension reaction. The CC cidentity of a SNP and for genotyping nucleic acid sample by CC cidentity of a SNP and for genotyping nucleic acid samples, for e.g. to cassess by association analysis the genotypic of an individual or group of caused by one or more SNPs. Phenotypic trait suspected of being CC agammaglobulinaemia, diabetes insipidus, Lesch-Nyhan syndrome, muscular dystrophy, familial hypercholesteroleemia, polycystic kidney disease, osteogenesis imperfecta and acute intermittent porphyria. Phenotypic craits also include symptoms of or susceptibility to multifactorial cdisease, including, rheumatoid arthritis, multiple sclerosis, inflammation, cancer, nervous system diseases and infection by pathogenic microorganism. The method is also useful in forensic investigations and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Single nucleotide polymorphism; SNP; single nucleotide primer extension;
SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SNP specific SNPE primer SEQ
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                                                                                                                                                                                                                                                                                                                                                                                               Claim 1; Page
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15-OCT-1999;
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Pred.
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No. 1.
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RESULT 454
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Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide cc primer extension (SNPE) primers, and the sequences of regions flanking clies of single nucleotide polymorphisms SNPs. The present invention cc includes kits for determining the presence or absence of a SNP, using the cligonucleotides of the invention. The PCR primers are used to amplify a CC SNP flanking sequence, the SNPE primer is used as a genotyping primer. CC cligonucleotides are useful for genotyping a nucleic acid sample by cc performing a single-nucleotide primer extension reaction. The CC cligonucleotides are useful for determining the presence, absence or cc identity of a SNP and for genotyping nucleic acid sample by cc assess by association analysis the genotype of an individual or group of cc individuals, having a pathological phenotypic trait suspected of being cagement by one or more SNPs. Phenotypic traits include diseases e.g. costeogenesis imperfecta and acute intermittent porphyria. Phenotypic traits also include symptoms of or susceptibility to multifactorial cd disease of which a component is or may be genetic such as autoimmune cc diseases, including, rheumatoid arthritis, multiple sclerosis,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 25
                                                                                                                                                                                                                                                                                                                                                                    acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer;
                                                                                                                                                                                                                                                                                                                                                                                              New genotyping oligonucleotide, useful for detecting the presence,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14-AUG-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13-OCT-2000; 2000WO-US028436
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Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia;
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C;
                                                                                                                                                                                                                                                                                                                                                                                      single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25
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Pred. No. 1
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Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide CC primer extension (SNPE) primers, and the sequences of regions flanking CC sites of single nucleotide polymorphisms SNPs. The present invention CC includes kits for determining the presence or absence of a SNP, using the CC oligonucleotides of the invention. The PCR primers are used to amplify a CC SNP flanking sequence, the SNPE primer is used as a genotyping primer. CC The oligonucleotides are useful for genotyping a nucleic acid sample by performing a single-nucleotide primer extension reaction. The CC oligonucleotides are useful for determining the presence, absence or CC identity of a SNP and for genotyping nucleic acid sample by CC assess by association analysis the genotype of an individual or group of CC individuals, having a pathological phenotypic trait suspected of being CC caused by one or more SNPs. Phenotypic trait suspected of being CC agammaglobulinaemia, diabetes insipidus, Lesch-Nyhan syndrome, muscular CC dystrophy, familial hypercholesterolaemia, polycystic kidney disease, CC osteogenesis imperfecta and acute intermittent porphyria. Phenotypic traits also include symptoms of or susceptibility to multifactorial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
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Matches 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nucleic acid sample.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis; inflammation; forensic investigation; paternity analysis; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 1; Page 65; 83pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              13-OCT-2000; 2000WO-US028436.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ7; MDZ12; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss.
                                                                                                                                                                                                                  New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ4, MDZ7 or MDZ12, e.g. cancer.
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Pred. No. 1.3e+03;
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proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is encoded at chromosome 7q22.1, MDZ4 is encoded at chromosome 1p21.3-22.2, MDZ7 is encoded at chromosome 1p21.2 and MDZ12 is encoded at chromosome 1sq26.1. The MDZ3, MDZ4, MDZ7, and MDZ12 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12. The probes are alterations in MDZ3, MDZ4, MDZ7, or MDZ12 genetic Locus. The probes are

The present invention relates to novel human zinc finger-containing

useful in constructing microarrays for measuring gene expression. proteins are useful as therapeutic agents for gene therapy or as vaccines. The present sequence was used to illustrate the inventic

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RESULT 457
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                                                                                                                                                          proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is encoded at chromosome 7g22.1, MDZ4 is encoded at chromosome 6p21.3-22.2, MDZ7 is encoded at chromosome 16p11.2 and MDZ12 is encoded at chromosome 12g26.1. The MDZ3, MDZ4, MDZ7, and MDZ12 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12, Gross are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ7; MDZ12; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADB04614 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                       New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ7 or MDZ12, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           02-AUG-2001; 2001US-00922181
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human MDZ7 scanning oligonucleotide SEQ ID 5600.
                                                                                                                             useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as
                                                                                                                                                                                                                                                                                                                             The present invention relates to novel human zinc finger-containing
                                                                                                                                                                                                                                                                                                                                                            Example 8; SEQ ID NO 5600; 103pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2003-423107/40
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                            Similarity
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                                                                                                                   The present sequence was used to illustrate the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gu Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP; 3 A;
                                                                                       BP; 4 A; 7 C; 10 G; 4 T; 0 U; 0 Other;
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                             Conservative
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                                           2.0%;
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                            Score 20.2; D
Pred. No. 1.3e
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                                           .3e+03;
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AXU

ADB04684;

ADB04684 RESULT 459

ADB04684 standard; DNA;

25 ВP

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RESULT 458
ADB04577
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AC ADB045
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                                                                                                                                                                                                                                                                                                                          The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is encoded at chromosome 702.1, MDZ4 is encoded at chromosome 722.1, MDZ4 is encoded at chromosome 722.1, MDZ4 is encoded at chromosome fp21.3-22.2, CMDZ7 is encoded at chromosome 16p11.2 and MDZ12 is encoded at chromosome 15p26.1. The MDZ3, MDZ4, MDZ7, and MDZ12 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder cossociated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic caids can also be used as probes to detect and characterize gross are useful in constructing microarrays for measuring gene expression. The probes are useful in constructing microarrays for measuring gene expression. The protein are useful as therapeutic agents for gene therapy or as caccines. The present sequence was used to illustrate the invention.
                                                                                                                             Matches
                                                                                                                                                             Query Match
Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human MDZ7 scanning oligonucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   02-AUG-2001; 2001US-00922181
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                                                                                                                                                                                                                                                              Sequence 25
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                                                           TATTTTAATTTTTGAGACAGAGT 629
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               or MDZ12, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gu Y,
                                                                                                                                                                                                                                                           BP; 4 A; 1 C; 4 G; 16 T; 0 U; 0 Other;
                                                                                                                             Conservative
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Pred. No. 1.
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RESULT 460
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Best Local
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Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ7; MDZ12; chromosome 7 chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ4, MDZ7 or MDZ12, e.g. cancer.
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                                                                                                                         Human MDZ7 scanning oligonucleotide SEQ ID 5669.
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                                                                                                                                                                                                                                                   ADB04683;
                                                                                                                                                                                                                                                                                                                ADB04683 standard;
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Pred. No. 1.
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   chromosome 15q26.1; cancer;
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The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is encoded at chromosome 7g22.1, MDZ4 is encoded at chromosome 6p21.3-22.2, MDZ7 is encoded at chromosome 16p11.2 and MDZ12 is encoded at chromosome 15p26.1. The MDZ3, MDZ4, MDZ7, and MDZ12 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12 genetic locus. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as
                                                                                                                                                                                                                                                                                                                                                                                                                                       New zinc finger-containing proteins and nucleic acids, useful manufacturing a medicament for treating or preventing a disorcassociated with decreased or increased expression or activity
                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                  Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Shannon M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                02-AUG-2001; 2001US-00922181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30-JUL-2002; 2002EP-00016874
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       developmental disorder;
                                                                              vaccines.
                                                                                                                                                                                                                                                                                                                                                                                                                   MDZ4, MDZ7 or MDZ12, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                  8; SEQ ID NO 5669;
                                       25 BP; 3 A; 9 C; 6 G; 7 T; 0 U; 0 Other;
                                                                            The present sequence was used to illustrate the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gu Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nguyen C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           88.
                                                                                                                                                                                                                                                                                                                                                                                103pp; English.
   В
 ۲.
Length
                                                                                                                                                                                                                                                                                                                                                                                                                                           disorder
tivity of MDZ3,
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RESULT 461 밁 S Query Match Best.Local S Matches 22 Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ7; MDZ12; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; Human MDZ7 ADB04576 developmental 537 \vdash l Similarity 22; Conserv standard; DNA; 25 CCTGCCTCAGCCTCCCAAGTAGCTG scanning oligonucleotide SEQ ID 2.0%; nilarity 88.0%; Conservative (first entry) 0 Score 20.2; Pred. No. 1 Mismatches 561 25 1.3e+03ω --Indels 0 Gaps 0,

EP1281758-A2 Homo sapiens.

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RESULT 462
ADB04681
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Best Local S
Matches 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is encoded at chromosome 7q22.1, MDZ4 is encoded at chromosome 6p21.3-22.2, MDZ7 is encoded at chromosome 16p1.2 and MDZ12 is encoded at chromosome 15q26.1. The MDZ3, MDZ4, MDZ7, and MDZ12 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12 genetic locus. The probes are alterations in MDZ3, MDZ4, MDZ7, or MDZ12 genetic focus. The probes are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ7 or MDZ12, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Shannon M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30-JUL-2002;
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                                                                                                                                                                                                                                   Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ7; MDZ12; Chromosome 7g22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              useful in constructing microarrays for measuring gene expression. proteins are useful as therapeutic agents for gene therapy or as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example
                                                                                                                                                                       EP1281758-A2
                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                             Human MDZ7 scanning oligonucleotide SEQ ID 5667.
                                                                                                                                                                                                                                                                                                                                                 20-NOV-2003
                                                                                                                                                                                                                                                                                                                                                                                                              ADB04681 standard; DNA;
                                                                                                           30-JUL-2002; 2002EP-00016874
                                                                           02-AUG-2001; 2001US-00922181
                                              (AEOM-) AEOMICA INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2003-423107/40
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22; Conserv
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llarity 88.0%;
Conservative
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                Nguyen
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Pred. No. 1
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XXFFFFFX 8X3030303030303030XXX
                                     CC proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is CC encoded at chromosome 7q22.1, MDZ4 is encoded at chromosome 6p21.3-22.2, CC MDZ7 is encoded at chromosome 16p11.2 and MDZ12 is encoded at chromosome 6p21.3-22.2, CC mDZ7 is encoded at chromosome 6p21.3-22.2, CC mDZ3, MDZ4, MDZ7, and MDZ12 sequences are useful in therapy, CC associated with decreased or increased expression or activity of MDZ3, CC MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic CC acids and proteins are also useful for diagnosing or monitoring a disease CC caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic CC acids can also be used as probes to detect and characterize gross CC alterations in MDZ3, MDZ4, MDZ7, or MDZ12 genetic locus. The probes are CC useful in constructing microarrays for measuring gene expression. The CC proteins are useful as therapeutic agents for gene therapy or as CC vaccines. The present sequence was used to illustrate the invention.
                                                                                                                                                                                                                                                                                                                                                                                          The present invention relates to novel human proteins and their coding sequences: MDZ3, MD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New zinc finger-containing proteins and nucleic acids, useful manufacturing a medicament for treating or preventing a disorcassociated with decreased or increased expression or activity MDZ4, MDZ7 or MDZ12, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 8; SEQ ID NO 5667; 103pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                           zinc finger-containing
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MDZ3,
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á Query Match Best Local S Matches 22 535 CTCCTGCCTCAGCCTCCCAAGTAGC Similarity Conservative 2.0%; <u>,</u> Score 20.2; Pred. No. Mismatches 559 1.3e+03 DB 1; Length 25; Indels 0, Gaps

0

Sequence 25

BP; 3 A; 10 C; 5 G; 7 T; 0 U; 0 Other;

RESULT 463 ADP70378/c ADP70378; ADP70378 standard; DNA; 25 ₽P

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CTCCTGCTTCAGTCTCCCGAGTAGC

25

Probe used to anlyse human testin-related gene (TRG) expression.

12-AUG-2004

(first entry)

human leukocyte antigen; HLA-B52; HLA-B62; cytotoxic T-cell; CTL; TRG2-41; TRG1-20; cytostatic; epithelial cancer; lung; stomach; c prostate; melanoma; vaccine; human; testin-related gene; ss; prob colon;

Homo sapiens.

JP2004141154-A

20-MAY-2004.

29-SEP-2003; 2003JP-00338402

30-SEP-2002; 2002JP-00286676

(/YOTI)

WPI; 2004-382710/36.

Novel tumor antigens TRG1-20 and TRG2-41 capable of recognizing and inducing human leukocyte antigen B52 or B62 constraint property of cytotoxic T lymphocyte, useful for treating cancer e.g., colon cance prostatic cancer, cancer,

SEQ IJ ö 9 34pp; Japanese

The invention relates to a capable of recognising and capable novel peptide comprising a TRG1-20 sequence inducing the human leukocyte antigen (HLA). (HLA) -B52

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RESULT 464
ANT73704
ID ANT737
AC ANT
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Best Local S
Matches 22
                                                                                               PCR primers AAT73703-4 were used to prepare probes (containing Alu repeats) for detecting a mutation in the locus of chromosome 14 associated with a presentle form of Alzheimer's disease. Each of the probes hybridises with one of the two human chromosomal DNA segments cloned in the CEPH yeast artificial chromosome (YAC) library under the accession numbers YAC 934A3 identifiable by genetic marker D14576). The probes are useful for diagnosis of the form of Alzheimer's disease associated with chromosome 14 by a method comprising making a preparation of metaphase chromosome 14 by a method comprising making a preparation conditions with the pair of probes or with one of the probes and another probe that hybridised probes and their relative positions on a significant number of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      or HLA-B62 constraint property of a cytotoxic T-cell (CTL) or a peptide comprising a TRG2-41 sequence capable of recognising and inducing the HI B52 of a CTL. The peptide of the invention demonstrates cytostatic activity and may be useful for inducing a cytotoxic T-cell in order to treat cancer, preferably epithelial cancer, more preferably lung cancer, stomach cancer, colon cancer, prostatic cancer and/or melanoma. The treatment may comprise the use of a vaccine. The current sequence is the of the probe of the invention which was used to analyse human testin-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 1; Page 8; 21pp; French
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Probes for diagnosing Alzheimer's disease - hybridising with chromosome 14 segments cloned in yeast artificial chromosome library.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1997-353201/33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       28-OCT-1994;
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22; Conserv
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Sequence

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BP; 4 A;

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4 T;

0 U;

0 Other

RESULT 466 AAV85582 ID AAV855

AAV85582 standard; DNA; 20

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RESULT 465
AAT73703/c
ID AAT737
XX AAT737
XX PCR pr
XX PR2742
PD 27-JUN
XX 28-OCT
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                                                                                                                                                                                                                                                                          repeats) for detecting a mutation in the locus of chromosome 14 associated with a presentle form of Alzheimer's disease. Each of the probes hybridises with one of the two human chromosomal DNA segments cloned in the CEPH yeast artificial chromosome (YAC) library under the accession numbers YAC 934A3 identifiable by genetic marker D14S251) and YAC 854F5 (identifiable by genetic marker D14S251) and chromosome 14 by a method comprising making a preparation of metaphase chromosome 14 by a method comprising making a preparation of metaphase chromosome from the patient's lymphoblastoid cells on a microscope slide, contacting the preparation under DNA hybridised make probe that the pair of probes or with one of the probes and another probe that hybridised probes and their relative positions on a significant number of pairs of chromosome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAT73703 standard;
                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PCR primers AAT73703-4 were used to prepare probes (containing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Probes for diagnosing Alzheimer's disease - hybridising with chromosome 14 segments cloned in yeast artificial chromosome library.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Weissenbach J, Heilig
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YAC; chromosome 14; presenile;
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RESULT 467
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LRP5
                                                                                                                                                                                                                                                                                  The present invention describes LRP5 (low density lipoprotein (LDL) receptor related protein, previously designated LRP-3). AAV85552 to AAV85556 represent PCR primer used for obtaining LRP5 cDNA. Nucleic acid molecules (NAMs) encoding LRP5 can be used for determining if an individual is susceptible to insulin dependent diabetes mellitus (IDDM). The NAMs or proteins can be used for reducing triglyceride levels in the serum of an individual. Therapies that affect LRP5 may also be useful in the treatment of autoimmune diseases such as glomerulonephritis, diseases and disorders involving disruption of endocytosis and/or antigen presentation, cytokine clearance and/or inflammation, viral infection, pathogenic bacterial toxin contamination, elevation of free fatty acids or hypercholesterolemia, type 2 diabetes, osteoporosis, Alzheimer's disease and cardiovascular disease. Products from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New isolated LDL-receptor related protein - used to develop products for treating, e.g. elevated triglyceride levels, diabetes, autoimmune disorders, inflammation or Alzheimer's disease.
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05-JUN-1997;
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                                                                                                                                                                                                                                             Sequence 20 BP; 4 A; 6 C; 5 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 12; Page 98; 200pp; English.
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P, Kawaguchi Y, Mer
lips MS, Twells RCJ;
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   (first entry)
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97US-0048740P.
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                                                            DNA;
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Merriman
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Metzker ML,
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Nakagawa
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Human mdm2 gene; proliferation; tumour; antisense; modulation; oligonucleotide; hyperproliferation; blood cancer; brain

07-JAN-2000 AAZ37736;

(first

entry)

mdm2

phosphorothioate

oligodeoxynucleotide #266.

phosphorothicate; p53; cancer; expression; inhibition; cancer; breast cancer;

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RESULT 468
AAZ37736/C
ID AAZ377
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DT 07-JAN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
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RESULT 469
AAZ37712/c
ID AAZ37712 standard; DNA; 20
XX
AC AAZ37712;
XX
DT 07-JAN-2000 (first entry)
XX
Human mdm2 phosphorothioat
XX
Human mdm2 gene; prolifera
KW Human mdm2 gene; prolifera
KW antisense; modulation; oli
KW hyperproliferation; blood
KW lung cancer; soft tissue c
XX
YN restenosis; ss.
XX
PN w09949065-A1.
XX
PN W09949065-A1.
XX
PP W09949065-A1.
XX
PP 30-SEP-1999.
XX
26-MAR-1999; 99WO-US0067
XX
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAZ37473-Z37738 represent human mdm2 phosphorothioate oligonucleotides. AAZ37471, AAZ37472, AAZ37739, AAZ37740 and AAZ37741 are used in the exemplification of the present invention. The present invention describes novel nucleotide antisense compounds, targetted to the 5' untranslated, translation termination codon, or 3' untranslated region of a nucleic acid encoding human mdm2, that modulates expression of human mdm2. The oligonucleotides mediate their effect by antisense inhibition of hyperproliferative gene expression. The antisense compound is used to treat an animal having a disease or condition associated with mdm2, particularly a hyperproliferative condition, more particularly cancer, especially of the blood, brain, breast, lung or soft tissue, or psoriasis, fibrosis, atherosclerosis or restenosis
                                                                                                                                                                            antisense; modulation; oligonucleotide; expression; inhibition; hyperproliferation; blood cancer; brain cancer; breast cancer; hung cancer; soft tissue cancer; psoriasis; fibrosis; atherosclerosis;
                                                                                                                                                                                                                                                             Human mdm2 phosphorothioate
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                     99WO-US006702.
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RESULT 470
RAZ3773/C
ID AAZ377/C
XX AAZ377
XX AAZ377
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DT 07-JAN
XX Human
XX Human
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAZ37473-Z37738 represent human mdm2 phosphorothioate oligonucleotides. AAZ37471, AAZ37472, AAZ37739, AAZ37740 and AAZ37741 are used in the exemplification of the present invention. The present invention describes novel nucleotide antisense compounds, targetted to the 5' untranslated, translation termination codon, or 3' untranslated region of a nucleic acid encoding human mdm2, that modulates expression of human mdm2. The oligonucleotides mediate their effect by antisense inhibition of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human mdm2 gene; proliferation; tumour; phosphorothioate; p53; antisense; modulation; oligonucleotide; expression; inhibition; hyperproliferation; blood cancer; brain cancer; breast cancer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human mdm2 phosphorothioate oligodeoxynucleotide
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Pred. No.
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Example

9; Page 55; 157pp; English

New antisense compounds used to treat

eg.

hyperproliferative conditions

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RESULT 471
AAA96410/c
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Autoimmune disease; polymorphic microsatellite repeat; PMR; CD28 gene; CTLA4 gene; costimulatory receptor gene locus; CGRL; lupus; insulin-dependent diabetes mellitus; IDDM; Addison's disease; leprosy; Graves disease; autoimmune hypothyroidism; myasthenia gravis; thymoma; thyroiditis; postpartum thyroiditis; rheumatoid arthritis; the postpartum thyroiditis; rheumatoid arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Primer used
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                                                                                    PCR primers AAA96409-10 were used to amplify polymorphic microsatellite repeat (PMR) sequences from the human costimulatory receptor gene locus (hCGRI). The primers are used in the method of the invention. The specification describes a method for determining the predisposition of a human subject to develop autoimmune disease. The method comprises detecting a PMR sequence in the CD28, ICOS gene or CTLA4 gene of the human costimulatory receptor gene locus (hCGRI). PMR sequences vary in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hashimoto's disease; coeliac disease; PCR primer;
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human costimulatory receptor gene locus (hCGRL). PMR sequences vary length among individuals and can be amplified to generate products differ in size. These products can then be detected by rapid and convenient high resolution processes. The method is useful for
                                                                                                                                                                                                                                                                                                                                                                                                                      Determining predisposition of humans to develop autoimmune disease involves detecting polymorphic microsatellite repeat sequence with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ling V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25-MAR-1999;
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                                                                                                                                                                                                                                                                                                                              Claim 18; Page 154; 160pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             to amplify a sara43/44 polymorphic microsatellite repeat.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INST INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                  polymorphic microsatellite
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Matches 20
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                                               The present sequence is that of interspersed repeated sequence PCR (I PCR) primer ALUS used to identify human-specific sequences in yeast artificial chromosomes (YAC) derived from the human chromosome 1q23.3 region. The chromosomal region contains the locus associated with absorptive hypercalciuria (AH). IRS-PCR fingerprints were generated, genes contained within YACs were identified by exon trapping. cDNA corresponding to the AH gene was isolated (see AAZS376). Identificat of the AH genomic region allows genetic screening for increased risk developing AH or osteoporosis with hypercalciuria
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; absorptive hypercalciuria; osteoporosis; nephrolithiasis; osteopathic; anticalciuric; chromosome 1g23.3-g24; therapy; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAZ35378 standard; DNA; 20
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                                                                                                                                                                                                                                                                                                                                                                                                     osteoporosis with
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hes 20; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                               genomic region useful in screening for absorptive hypercalciuria
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                                                                                                                                                                                                                                                                                                                                            Page 125; 153pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PCR primers AAA14945-47 were used to amplify DNA containing the repeated DNA sequences Alu. The amplified fragment was used as a probe of the invention. The specification describes probes which are used for chromosomal labelling. The probes consist of a set of DNA segments represented at higher level in certain chromosomal bands and produced by TRS-PCR (interspersed repeat sequence-polymerase chain reaction) using primers specific for the repeated DNA sequences Alu and LINE. The probes are used for studying karyotypes, including those associated with chromosomal rearrangements, particularly in multicolour fluorescent in
Human; glycogen synthase kinase 3 alpha; antidiabetic; cytostatic; antisense therapy; diabetes; hyperproliferative disorder; inflamma neurological disorder; tumour; haematopoietic disorder; infection;
                                                                                                      Human glycogen synthase kinase 3 alpha antisense oligo ISIS #116649.
                                                                                                                                                           01-NOV-2001
                                                                                                                                                                                                               AAD14808;
                                                                                                                                                                                                                                                            AAD14808 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Probes for chromosomal labeling, useful for diagnostic determination karyotype, are prepared by amplification using primers specific for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2000-318009/27
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                                                                                                                                                                                                                                                               DNA;
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The invention relates to an antisense compound 8 to 30 nucleobases in length targetted to a nucleic acid encoding Jyycogen synthase kinase 3 alpha. The antisense compound specifically hybridises with and inhibits the expression of glycogen synthase kinase 3 alpha. The antisense compound is useful for the treatment of a diseases associated with glycogen synthase kinase 3 alpha such as diabetes, a neurological disorder, a haematopoietic disorder, a hyperproliferative disorder or a developmental disorder. The antisense compounds may also be used prophylactically to prevent or delay infection, inflammation or tumour formation. The present sequence is a phosphorothicate antisense oligonucleotide targetted to human glycogen synthase kinase 3 alpha menemic nua
                     genomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              hyperproliferative disorder; developmental disorder; antisense; phosphorothioate backbone; ss.
                                                                                                                                                                       Example 15; Page 83; 115pp; English.
                                                                                                                                                                                                         Antisense compound 8 to 30 nucleobases in length comprising a that is targeted to a nucleic acid molecule encoding glycogen kinase 3 alpha, useful for the treatment of e.g. diabetes and
                                                                                                                                                                                                                                                        WPI; 2001-442247/47.
                                                                                                                                                                                                                                                                                                                           21-JAN-2000; 2000US-00488856
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== "Methoxyethyl residues"
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                                                                                                                                                                                                                                                                               Butler MM,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           residues"
                                                                                                                                                                                                                                compound
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Query Match Best Local Similarity

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Score 20; DB 1; Li Pred. No. 1.1e+03;

Length

Sequence

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Best Local Similarity
Matches 20; Conserv
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11-JAN-2000;
02-MAY-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to primers for synthesising full length cDNA clones. 830 cDNA molecules encoding a human protein have been isolated and nucleotide sequences of 5'- and 3'-ends of the cDNA molecules have been determined. Primers for synthesising the full length cDNA are useful for clarifying the function of the protein encoded by the cDNA. The full length clones were obtained by construction of full length enriched cDNA
                                                                                                                                                                                                                                                                                                                                              libraries that were synthesised by the oligo-capping method. The primers enable the production of the full length cDNA easily without any special methods. The present sequence is a primer used to amplify a human cDNA clone provided in the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EP1130094-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          830 Primers useful for synthesizing full length cDNA clones and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Wakamatsu
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           07-JUL-2000; 2000EP-00114089
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                                                                                                                                                                                                                                                                                                               Sequence
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                     02-MAY-2001
                                                        AAF80866;
                                                                                      AAF80866 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            genetic
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                                                                                                                                                                                                                                                                                                             BP; 4 A;
                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      99JP-00194486.
2000JP-00118774.
2000JP-00183765.
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                     (first entry)
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                                                                                                                                                                                                                                                                                                               7 C;
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a T, Nagai
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                                                                                                                                                                                                                                                                                                               3 G; 6 T; 0 U;
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K, Kojima
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                                                                                                                                                                                                                                                            DB 1; Le . 1.1e+03;
                                                                                                                                                                                                                                                                                                                 0 Other;
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T, Koga
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AAF80891/c
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Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to an antisense compound 8-30 nucleobases in length targeted to nucleobases 1-308 of the 5' untranslated region, 1776-1806 of the translation termination codon region or 1818-2370 of the 3' untranslated region of a nucleic acid molecule encoding human mdm-2. The invention is useful for reducing hyperproliferation of human cells, modulating the expression of mdm2 in human cells or tissues or in vitro. The hyperproliferative disorder includes cancer or psoriasis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel antisense compound 8-30 nucleobases in length targeted to a nucleic acid molecule encoding human mdm-2 useful for modulating the expression of human mdm-2 and reducing hyperproliferation of human cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human mdm2 phosphorothioate oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2001-190948/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 06-FEB-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26-MAR-1998;
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                                                                                                                                                               26-MAR-1999;
                                                                                                                                                                                               06-FEB-2001.
                                                                                                                                                                                                                                                                  Homo
                                                                                                                                                                                                                                                                                                  Antisense; mdm2;
                                                                                                                                                                                                                                                                                                                                     Human mdm2
                                                                                                                                                                                                                                                                                                                                                                                                           AAF80891;
                                                                                                                                                                                                                                                                                                                                                                                                                                            AAF80891 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (ISIS-) ISIS
                         WPI; 2001-190948/19
                                                         Miraglia LJ,
                                                                                                                              26-MAR-1998;
                                                                                                                                                                                                                                US6184212-B1
                                                                                            (ISIS-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 937 CTGTTACCCAGGCTGGAGTG
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                                                                                            ISIS
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                                                                                                                                                                                                                                                                                                                                     phosphorothioate oligonucleotide #265.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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                                                                                            PHARM INC
                                                                                                                                                                                                                                                                                                                                                                         (first entry)
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                                                         Nero P,
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                                                                                                                            98US-00048810
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                                                                                                                                                                                                                                                                                                  hyperproliferation; cancer; psoriasis;
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100.0%; Pr
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                                                                                                                                                                                                                 The present invention relates to an antisense compound 8-30 nucleobases in length targeted to nucleobases 1-308 of the 5' untranslated region, 1776-1806 of the translation termination codon region or 1818-2370 of the 3' untranslated region of a nucleic acid molecule encoding human mdm-2. The invention is useful for reducing hyperproliferation of human cells, modulating the expression of mdm2 in human cells or tissues or in vitro.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel antisense compound 8-30 nucleobases in length targeted to a nucleic acid molecule encoding human mdm-2 useful for modulating the expression
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Miraglia LJ,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (ISIS-) ISIS
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                                                                                                                                                                                             hyperproliferative disorder includes cancer or psoriasis
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  hyperproliferation; cancer; psoriasis; ss.
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                             2.0%;
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                                                                                                                           C; 6 G; 4 T; 0 U; 0 Other;
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                             Score 20; DB 1;
Pred. No. 1.1e+0
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1.1e+03;
                                 1.1e+03;
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                                                               Length 20;
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Score 20; Pred. No.

DB 1; 1.1e+03

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                                                                                   CC includes kits for determining the presence of a SNP, using the CC oligonucleotides of the invention. The PCR primers are used to amplify a CC SNP flanking sequence, the SNPs primer is used as a genotyping primer. CC The oligonucleotides are useful for genotyping a nucleic acid sample by CC performing a single-nucleotide primer extension reaction. The CC oligonucleotides are useful for determining the presence, absence or CC oligonucleotides are useful for determining the presence, absence or CC identity of a SNP and for genotyping nucleic acid samples, for e.g. to CC assess by association analysis the genotype of an individual or group of CC individuals, having a pathological phenotypic trait suspected of being CC caused by one or more SNPs. Phenotypic traits include diseases e.g. CC agammaglobulinaemia, diabete includent traits include diseases, cost of traits also include symptoms of or susceptibility to multifactorial CC diseases, including, rheumatoid arthritis, multiple sclerosis, pathogenic confiammation, cancer, nervous system diseases and infection by pathogenic confiammation, cancer, nervous system diseases and infection by pathogenic confiammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmuno diacute intermittent porphyria.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SNP
                                                                                                                                                                                                                                                                                                                                                                                                                               primer extension (SNPE) primers, and the sequences of regions flanking sites of single nucleotide polymorphisms SNPs. The present invention
                                                                         inflammation, cancer, nervous system diseases and infection by pathogenic microorganism. The method is also useful in forensic investigations and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1; Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Picoult-Newburg L,
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                                                       paternity analysis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      acid sample.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ystic kidney disease; osteogenesis imperfecta; autoimmune disease; intermittent porphyria; rheumatoid arthritis; multiple sclerosis; mation; forensic investigation; paternity analysis; PCR primer; s
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                                  containing
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                                                   The present sequence represents
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The present invention relates to antisense compounds, 8-30 nucleobáses in CC length targeted to the 5' untranslated region, translation termination CC codon region, 3' untranslated region, coding region or translation start CC site of a nucleic acid encoding human mdm2, where the antisense compound CC modulates the expression of human mdm2. The antisense oligonucleotides of CC the invention are useful for encoding human mdm2 and for inhibiting the expression of human mdm2. They may be used for treating an animal having CC adisease or condition associated with amplification of mdm2 gene or CC expression of mdm2 e.g. a hyperproliferative disorder such as cancer CC (blood, brain, breast, lung, or a soft tissue cancer) and psoriasis, CC fibrosis, atherosclerosis or restenosis, tumours, colorectal carcinoma and chronic myelogenous leukemia. The antisense compound may be CC administered with a chemotherapeutic agent to overcome drug resistance. The antisense compound, is also useful CC method, which involves the use of the antisense compound, is also useful CC method, which involves the use of the antisense compound, is also useful CC diagnostic tools. AAS29242-AAS29507 represent the human mdm2 antisense
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26-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 9; Page 18; 81pp; English.
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(MONI/)
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sclerosis; tumour; cytostatic; anti psoriatic;
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               The present invention relates to antisense compounds, 8-30 nucleobases in length targeted to the 5' untranslated region, translation termination codon region, 3' untranslated region, coding region or translation start site of a nucleic acid encoding human mdm2, where the antisense compound modulates the expression of human mdm2. The antisense oligonucleotides of the invention are useful for encoding human mdm2 and for inhibiting the expression of human mdm2. They may be used for treating an animal having a disease or condition associated with amplification of mdm2 gene or overexpression of mdm2 e.g. a hyperproliferative disorder such as cancer (blood, brain, breast, lung, or a soft tissue cancer) and psoriasis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; mdm2; hyperproliferative disorder; cancer; psoriasis;
atherosclerosis; tumour; cytostatic; anti psoriatic;
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26-MAR-1999;
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) GRAHAM M J.
) MONIA B P.
) COWSERT L M.
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1-6 and bases 15-20 are 2'-0-
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              The present invention relates to antisense compounds, 8-30 nucleobases length targeted to the 5' untranslated region, translation termination codon region, 3' untranslated region, coding region or translation stan
                                                                                            nucleobases targeted a region (e.g. translation of a nucleic acid encoding human mdm2.
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26-MAR-1999;
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CC modulates the expression of human mdm2. The antisense oligonucleotides of CC the invention are useful for encoding human mdm2 and for inhibiting the CC expression of human mdm2. They may be used for treating an animal having CC a disease or condition associated with amplification of mdm2 gene or CC (blood, brain, breast, lung, or a soft tissue cancer) and psoriasis, CC fibrosis, atherosal-erosis or restenosis, timeours, colorectal carcinoma CC and chronic myelogenous leukemia. The antisense compound may be administered with a chemotherapeutic agent to overcome drug resistance. CC administered with a chemotherapeutic agent to overcome drug resistance. CC The antisense compound reduces hyperproliferation of human cells. The CC method, which involves the use of the antisense compound, is also useful CC for detecting the role of mdm2 expression in various cell functions and CC diagnostic tools. AAS29242-AAS29507 represent the human mdm2 antisense CC oligonucleotides of the present invention
  Sequence
  20
  B₽;
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7 C;
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Query Match Best Local S Matches 20 Similarity 20; Conserv Conservative 2.0%; 0; Score 20; Pred. No. Mismatches 1.1e+03; Indels ٥, Gaps

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937
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                   CTGTTACCCAGGCTGGAGTG 956
CTGTTACCCAGGCTGGAGTG
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AAK98932 ID AAV AAK98932 standard; DNA; 20

24-MAY-2002 (first entry)

Human Beta-globin 5' MAR antisense primer BMR1.

Expression vector; beta-globin nuclear matrix attachment region; MAR; SV40 virus; gastrin; tumour growth factor beta soluble receptor II; TGF-beta SRII; TGF-beta-overexpressed disease; human; PCR; primer; ss 88

Homo sapiens

WO200214525-A2

27-JUL-2001; 2001WO-KR001285

29-JUL-2000; 2000KR-00043996

(MOGA-) MOGAM BIOTECH LAB RES INST.

WTW Ç Kim J, 유 ıs, Yoon J, Baek Ķ Chung တ Park

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Yoon

Ķ,

WPI; 2002-269202/31.

New expression vectors for use in animal cells (e.g. pMS, pSG evectors), useful for producing recombinant proteins in various cells and recombinant protein having a unique structure and fur function. and and pMSG

Example 1; Page 77; 85pp; English

The invention relates to new expression vectors for animal cells comprising a beta-globin nuclear matrix attachment region (MAR) sequence or its complementary sequence at 5'-terminal end of a promoter and/or a SV40 virus poly-A signal and transcription termination site of gastrin gene. The expression vectors are useful for producing recombinant proteins in various animals cells and recombinant protein having a unique structure and function. The vectors, which have increased expression efficiency and levels for foreign genes, are useful for expressing foreign proteins used in an animal cell system, e.g. tumour growth factor

of a

nucleic

acid encoding

human mdm2,

where the

antisense

compound

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RESULT 484
ABS592C
ID ABS592
AC ABS692
AC ABS6
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; ss; antisense; cellular apoptosis susceptibility gene; antiinflammatory; antitumour; cytostatic; CAS; CSE1; CSP; chromosome 20q13; mitosis; apoptosis; proliferation; cancer; importin-alpha; nuclear localisation; cell cycle; hyperproliferative disorder; degenerative disorder; Alzheimer's disease; parkinson's disease; amyotrophic lateral sclerosis; ALS; retinitis pigmentosa; blood cell disorder; gene therapy; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human CAS
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                                                                                               New antisense compound that hybridizes and inhibits nucleic acid encoding cellular apoptosis susceptibility gene, useful for treating a hyperproliferative disorder such as cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        modified_base
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                                                                                                                                                                                                                                                   Cowsert LM,
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                                                      Claim 3; Page
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                                                                                                                                                                                                                                                   Freier SM;
                                                                                                                                                                                                                                                                                                    PHARM INC.
                                                      91; 135pp; English.
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16. .20
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Pred. No.
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The invention discloses

antisense compounds,

of,

8

50 nucleobases

New antisense oligonucleotides targeted to nucleic acid encoding casein kinase 2-alpha prime, useful for diagnosing and/or treating a disease of condition associated with expression of casein kinase 2-alpha prime.

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Claim 3; Page 94; 129pp; English

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ABS67840/
ID ABS6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    밁
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Best Local Similarity
Matches 20; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; casein kinase 2-alpha prime; diabetes mellitus; hyperproliferative disorder; breast cancer; prostate cancer; liver cancer; infection; inflammation; tumour formation; cytostatic; antidiabetic; antiinflammatory; antimicrobial; phosphorothioate;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              antisense therapy;
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                                                                                                                                                                                                                                                                                                                                                                                                               08-FEB-2001; 2001US-00780173
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Pred.
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RESULT 486
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therapeutically or prophylactically to treat an animal having a disease or condition associated with caspase 6, such as Rieger's syndrome or ataxia telangiectasia, hyperproliferative disorder, a hamatopoietic disorder, a bone metabolism or cholesterol disorder, various types of
                                                                                                    nucleotides in length that is targeted to a nucleic acid molecule encoding caspase 6, where the oligonucleotide specifically hybridises with and inhibits the expression of caspase 6. The oligonucleotide of the invention specifically hybridises to and inhibits expression of caspase in cells or tissues. The oligonucleotides can be administered
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                                                                                                                                                                                                                                                                                               Example 15; Page 89; 141pp;
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AAL40354
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맑
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity Matches 20; Conserv
                                                         Matches
                                                                     Query Match
Best Local
                                                                                                                                                    The invention relates to an antisense oligonucleotide compound of 8 to 50 nucleotides in length that is targeted to a nucleic acid molecule encoding caspase 6, where the oligonucleotide specifically hybridises with and inhibits the expression of caspase 6. The oligonucleotide of the invention specifically hybridises to and inhibits expression of caspase 6 in cells or tissues. The oligonucleotides can be administered therapeutically or prophylactically to treat an animal having a disease or condition associated with caspase 6, such as Rieger's syndrome or ataxia telangiectasia, hyperproliferative disorder, a hone metabolism or cholesterol disorder, various types of cancer, neurological conditions such as Alzheimer's disease and other deregulated apoptotic pathological conditions. This polynucleotide sequence represents a human caspase 6 oligonucleotide relating to the invention. NOTE: This phosphorothioate oligonucleotide sequence has 2'-MOE wings and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Muscular; cytostatic; nootropic; neuroprotective; ophthalmological; antilipaemic; osteopathic; caspase 6; Rieger's syndrome; bone metabolism; ataxia telangiectasia; hyperproliferative disorder; cholesterol disorder; haematopoletic disorder; cancer; neurological; Alzheimer's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human caspase 6 antisense inhibition related oligo SEQ ID No
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAL40354;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  04-OCT-2000; 2000US-00679299.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200229066-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   apoptotic; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Brown-Driver VL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               03-OCT-2001; 2001WO-US030871.
                                                                                                                                                                                                                                                                                                                                                                                    Claim
                                                                                                                                                                                                                                                                                                                                                                                                               An antisense oligonucleotide of 8 to 50 nucleotides in length that inhibits caspase 6, is useful for treating Rieger's syndrome.
                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            2002-471315/50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         387
                           211
                                                                                                                                                                                                                                                                                                                                                                                    3; Page
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                                                         20;
 _
                                                                                                                                             gap
                                                                     Similarity
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                                                                                                                 20
                       CTGGTCTCGAACTCCCGACC 230
 CTGGTCTCGAACTCCCGACC 20
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                                                                                                                 BP;
                                                       2.0%;
ilarity 100.0%;
Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                  89; 141pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA;
                                                                                                               9 C; 4 G; 4 T;
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                                                         Score 20; DB; Pred. No. 1.1
0; Mismatches
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                                                                                                                 0 U;
                                                                                   DB 1;
                                                                     1.1e+03;
                                                                                                                 0 Other;
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                                                                                   Length 20;
                                                            Indels
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RESULT 489

BXGXEXE

ABL44004;

11-APR-2002

(first entry)

Human chromosome 1p36-35 PCR primer SEQ ID NO:1048.

RESULT 490

ABL44004

ABL44004 standard; DNA;

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ABL44512/c
IID ABL445
XX H.
AC ABL445
XX JP2001
XX JP2001
XX JP2001
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XX IIAAA
PA (RIKA
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                                                                                                                                                               Query Match
Best Local S
                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human chromosome 1p36-35 PCR primer SEQ ID NO:1556
                                                                                                                                                                                                                                                                                                                                  reconstituted as the positions on the chromosome and arrayed. The microarray is useful for gene analysis. ABL42957 to ABL45322 represent PCR primers for human chromosome 1p36-35 DNA, and ABL43323 to ABL45634 represent PCR primers for human chromosome 21q22.1, which are specifically claimed for use in the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    JP2001321190-A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    11-APR-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 4; Page 35; 528pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2002-144136/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          12-MAR-2001; 2001JP-00068285
                                                                                                                                                                                                                                                                     Sequence 20 BP; 4 A; 4 C; 8 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10-MAR-2000; 2000JP-00066716.
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(GENO-) GENOTEX YG.
                                               542 CTCAGCCTCCCAAGTAGCTG 561
20
                                                                                                                                  20;
                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      genome clones.
CTCAGCCTCCCAAGTAGCTG 1
                                                                                                                                      Conservative
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                                                                                                                    2.0%; Pr/
100.0%; Pr/
0;
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                                                                                                                                                                      Score 20;
Pred. No.
                                                                                                                                         Mismatches
                                                                                                                                                                                                           DB 1;
                                                                                                                                                                      1.1e+03;
                                                                                                                                                                                                           Length 20;
                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               in the multiwell
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                                                                                                                                         Gaps
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RESULT 491
ABA92187
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local (
                                  Homo sapiens
                                                                     NALPN; nyctalopin; human; congenital stationary night blindness; CSNB; glycosylphosphatidylinositol; GPI; proteoglycan; retina; polymorphism; marker; PCR; primer; ss.
                                                                                                                                                                                      06-JUN-2002
                                                                                                                                                                                                                           ABA92187;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 4; Page 25; 528pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Arraying genome clones
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10-MAR-2000; 2000JP-00066716
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20-NOV-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; chromosome 1p36-35; chromosome 21q22.1; genetic analysis; genome;
                                                                                                                                               Polymorphism 506B13CA1 reverse PCR primer.
                                                                                                                                                                                                                                                                ABA92187 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12-MAR-2001; 2001JP-00068285
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                             384 CTCCCAAAGTGCTGGGATTA 403
                                                                                                                                                                                                                                                                                                                                                            L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RIKAGAKU KENKYUSHO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENOTEX YG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20
                                                                                                                                                                                                                                                                                                                                                        CTCCCAAAGTGCTGGGATTA 20
                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP; 5 A; 5 C; 5 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                 2.0%;
                                                                                                                                                                                                                                                                BP.
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Pred. No.
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385 TCCCAAAGTGCTGGGATTAC 404

Best Loc Matches Query Match

Local

l Similarity 20; Conserv

Conservative

0,

100.0%; 2.0%;

Score 20;

Pred. No. Mismatches

1.1e+03; DB 1; Length 20, 0

0

Gaps

0

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TCCCAAAGTGCTGGGATTAC

20

ABA92208 RESULT 492

ABA92208 standard; DNA; 20

ВP

Homo sapiens

14-MAY-2001; 2001CA-02345915

12-NOV-2001.

NYX; nyctalopin; human; congenital stationary night blindness; CSNB; glycosylphosphatidylinositol; GPI; retina; SLRP;

leucine-rich proteoglycan; therapy; diagnosis; PCR; primer; ss

Reverse PCR primer for polymorphism 506B13CA1.

06-JUN-2002

(first entry

ABA92208;

CA2306241-A1

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The present sequence is that of a reverse primer used, with the forward CC primer given in ABA92186, in PCR analysis of polymorphism 506B13CA1 CDX51042). This was 1 of 3 novel markers identified in a genotype CC analysis of x-linked congenital stationary night blindness (CSNB) CF families. The new, and some previously known, markers allowed an analysis of selected recombinant x chromosomes to determine the CSNB1 minimal CC physical map of the CSNB1 minimal region in xp11.4 was developed. This CC physical map of the CSNB1 minimal region in Xp11.4 was developed. This CC physical map of the CSNB1 minimal region in Xp11.4 was developed. This CC physical map of the CSNB1 minimal region in Xp11.5 An extended CC NALPN CDNA sequence (see ABA92185) was established by sequencing of PCR and RACE products obtained from retinal RNA. 11 Different mutations were CC identified in NALPN, none of which were observed in normal individuals. These included missense mutations, insertions and deletions of the coding region that are predicted to disrupt specific functions of nyctalopin, CC and may be informative as to the structure-function relationship of the CC and such informative as to the structure function relationship of the CC and such information of the NALPN gene will also provide a tool for the NALPN gene will also provide a tool for the control of the NALPN gene will also provide a tool for the control of the NALPN gene will also provide at tool for the control of the NALPN gene will also provide at tool for the control of the NALPN gene will also provide at tool for the control of the NALPN gene will also provide at tool for the control of the NALPN gene will also provide at tool for the control of the NALPN gene will also provide at tool for the control of the NALPN gene will also provide at tool for the control of the NALPN gene will also provide at tool for the control of the NALPN gene will also provide at tool for the control of the NALPN gene and the control of the NALPN gene and the control of the NALPN gene and th
Sequence 20 BP; 5 A; 5 C; 5 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 1; Page 28; 44pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel purified mammalian retinal, kidney glycosylphosphatidylinositol-
anchored small leucine-rich proteoglycan and polynucleotides encoding
them used to diagnose complete X-linked congenital stationary night
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12-NOV-2001
                                                                having this disorder
                                                                                                          the diagnosis of complete X-linked CSNB in individuals suspected of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               12-MAY-2000; 2000CA-02306241
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CThe present sequence is that of a reverse primer used, with the forward CC primer given in ABA92207, in PCR analysis of polymorphism 506B13CA1 (C (DXS10042). This was 1 of 3 novel markers identified in a genotype CC analysis of X-linked congenital stationary night blindness (CSNB) CC families. The new, and some previously known, markers allowed an analysis CC of selected recombinant X chromosomes to determine the CSNB1 minimal CC region. To identify candidate genes for the CSNB1 locus, a robust CC physical map of the CSNB1 minimal region in Xp11.4 was developed. This CC identified the NYX gene encoding nyctalopin (see AAM51131). An extended CC NYX CDNA sequence (see ABA92206) was established by sequencing of PCR and CC identified in NYX genes from retinal RNA. 14 Different mutations were CC identified in NYX genes from different CSNB families, none of which were observed in normal individuals. These included missense, insertion, stop and deletion mutations that are predicted to disrupt specific functions CC of nyctalopin. The invention provides a method and kit for diagnosing CC complete X-linked CSNB, which involves screening for alterations in the CC candidates that affect nyctalopin expression or production
                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 493
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel purified mammalian glycosyl-inositol phospholipid-anchored sm
leucine-rich proteoglycan, and genes encoding proteoglycan which are
useful for diagnosing complete X-linked congenital stationary night
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2002-242185/30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 5 A; 5 C; 5 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 1; Page 26; 65pp; English
                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                     cell growth
                                                                                                                                                                                                                                                                       Telomerase reverse transcriptase; TERT; cytostatic; apoptosis; cell growth inhibitor; antisense oligonucleotide; antisense technology;
                                                                                                                                                                                                                                                                                                                         Telomerase reverse transcriptase, antisense oligonucleotide #69
                                                                                                                                                                                                                                                                                                                                                          09-APR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                AAS96659 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (BECH/) BECH-HANSEN N T.
                                                16-MAY-2000;
07-DEC-2000;
                                                                                                                                                                     WO200188198-A1
                                                                                                 15-MAY-2001; 2001WO-US015774.
              (ISIS-) ISIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   385 TCCCAAAGTGCTGGGATTAC 404
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   l Similarity
20; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                 PHARM
                                                 2000US-00572423.
2000US-00733294.
                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                20
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     <u>.</u>
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863 TGCTGGGATTACAGGCGTGA 882

Query Match Best Local Matches

Similarity

100.0%;

2.0%;

20;

Conservative

0; Mismatches Score 20; Pred. No.

DB 1; Le 1.1e+03;

Length 20

0

Gaps

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transcriptase), where the compound specifically hybridises with and inhibits the expression of TERT. A series of oligonucleotides were designed to target different regions of the human TERT RNA. These were 20 nucleotides in length and composed of a central gap region consisting of ten 2'-deoxynucleotides, flanked on both sides (5' and 3' directions) by five-nucleotide wings. The wings were composed of 2'-methoxyethyl (2'-mothotides. The compounds were analysed for their effect on human TERT mRNA levels by reverse transcriptase (RT)-polymerase chain reaction (PCR). The compound is useful for inhibiting the expression of TERT in compound is useful for inhibiting the expression of TERT in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI;
                                               cells or tissues, for treating a human having disease or condition associated with TERT, for modulating apoptosis, for inhibiting cell growth (preferably, cancer cell growth), in antisense therapy and for diagnostics and therapeutics. This sequence is an antisense oligonuclectide used to modulate the activity of nucleic acid molecules encoding TERT, described in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New compound targeted to nucleic acid molecule encoding telomerase transcriptase (TERT), which specifically hybridizes with and inhibits expression of TERT, useful for modulating apoptosis and inhibiting cell
                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention describes a compound, 8-50 nucleobases in length targeted to a nucleic acid molecule encoding human TERT (telomerase reverse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Monia BP, Gaarde WA,
Sequence 20 BP; 4 A; 3 C; 8 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 19; Page 91; 154pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2002-075321/10.
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RESULT 494
ABK91100/c
TO PROPERTY OF STANDARD STANDA
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                                                                                                                                                                                                                                                                                 14-MAR-1990;
06-FEB-1992;
02-DEC-1993;
07-NOV-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human, fluorescent labelling technique, fluorescent intercalating nucleic acid detection; electrophoresis; diagnostic assay;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABK91100 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    05-DEC-2002
                                          WPI; 2002-722081/78
                                                                                                                          Glazer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10-OCT-2000; 2000US-00686147
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   06-AUG-2002.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABK91100;
                                                                                                                                                                                                      (REGC ) UNIV CALIFORNIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           labelling;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      'سر
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                                                                                                                          Mathies RA,
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                                                                                                                                                                                                                                                                                 90US-00493307.
92US-00831823.
93US-00161231.
97US-00966398.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    for human DNA derived from chromosome 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               primer; ss.
                                                                                                                              Peck K;
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention relates to novel fluorescent labelling techniques and fluorescent labels. The method and compositions of the invention are useful for detecting molecules using fluorescent labels where fluorescent intercalating dyes have strong non-covalently binding affinities for the dsDNA. The method is useful for detecting molecules e.g. nucleic acids, in electrophoresis methods. The method can also be applied to diagnostic assays and cell labelling. The fluorescent label is sensitive, stable and resistant to self-quenching. The present sequence represents a PCR primer used to amplify human DNA derived from chromosome 21
         WPI; 2003-184032/18
                                                                                                                                                                                                    modified_base
                                                                                                                                                                                                                                                                                                                                             Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                     Human; superoxide dismutase 1; antisense; neuroprotective; cytostatic; antiinflammatory; amyotrophic lateral sclerosis; apoptosis; hyperproliferative disorder; therapy; infection; inflammation; tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                      Human superoxide dismutase 1 antisense inhibitor #
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ACC40946 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Detection of separated molecules involves use of a group comprising double stranded DNA probe and fluorescent molecule.
                                Bennett FC,
                                                                           21-JUN-2001;
                                                                                                  19-JUN-2002; 2002WO-US019664
                                                                                                                          03-JAN-2003
                                                                                                                                               WO2003000707-A2
                                                                                                                                                                                                                                                 modified_base
                                                                                                                                                                                                                                                                                                         modified_base
                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
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                                                       (ISIS-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            388 CAAAGTGCTGGGATTACAGG 407
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20;
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                                                     SISI
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0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                               Dobie K;
                                                       PHARM INC
                                                                            2001US-00888360
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               8
                                                                                                                                                                                                      /note= '
                                                                                                                                                                                                                                                             methylcytosine"
                                                                                                                                                                               /*tag=
/mod_ba
                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
                                                                                                                                                                                                                            /*tag=
/mod_ba
                                                                                                                                                                                                                                                                         note= "Phosphorothicate linkages.
                                                                                                                                                                                                                                                                                     /mod
                                                                                                                                                                      note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%;
                                                                                                                                                                                                                                                                                     base=
                                                                                                                                                                                                                             base= OTHER
                                                                                                                                                                                  base= OTHER
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                                                                                                                                                                     "2'-methoxyethyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               English
                                                                                                                                                                                                                 -methoxyethyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ₽₽
                                                                                                                                                                                                                                                                                     OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
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Pred. No.
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                                                                                                                                                                      (2'-MOE) nucleotides"
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                                                                                                                                                                                                                 (EOM-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length
                                                                                                                                                                                                                  nucleotides"
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                                                                                                                                                                                                                                                                          A11
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Novel antisense compounds targeted to nucleic acids encoding human superoxide dismutase 1, for modulating expression of the dismutase treating diseases or conditions, e.g. amyotrophic lateral sclerosis 1, for modulating conditions, e.g. and

Example 15; Page 77; 107pp; English

The invention relates to a compound of 8-50 nucleobases in length, CC targeted to a nucleic acid molecule encoding human superoxide dismutase CC 1. The compound specifically hybridises with and inhibits the expression CC of human superoxide dismutase 1 by hybridising with at least an 8CC nucleobase portion of the nucleic acid molecule encoding the active site CC for the enzyme. The activity of compounds of the invention may be CC described as neuroprotective, cytostatic and antiinflammatory. The CC mechanism of action of compounds of the invention is antisense inhibition of human superoxide dismutase 1 expression by chimeric phosphorothicate CC oligonucleotides having 2'-methoxyethyl (2'-MOE) wings and a deoxy gap. CC compounds of the invention are useful for inhibiting the expression of human superoxide dismutase 1 in human cells or tissues, and for treating a disease or condition associated with this enzyme (antisense therapy), cespecially amyotrophic lateral sclerosis, a disease or condition arising CC used in diagnostics, therapeutics and as a research reagent, e.g. cused in diagnostics, therapeutics and as a research reagent, e.g. comparation. Sequences given in records ACC40880-ACC40957 represent human construction of tumour conditions are antisense inhibitor oligonucleotides

Sequence 20 BP; 5 A, 4 C; 8 ູດ 3 T; 0 u; o Other;

S Matches Query Match Local 729 Similarity AGTAGCTGGGACTACAGGCG 748 Conservative 2.0%; 0 Score 20; Pred. No. Mismatches DB 1; 1.1e+03; 0 Length 20; Indels 0 Gaps 0

RESULT 496 ABZ79385 ID ABZ793

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μ.

20

ABZ79385 standard; DNA; 20 ВP

ABZ79385

01-MAY-2003 (first entry

Acetyl-Coenzyme A-carboxylase-alpha gene PCR primer, SEQ Ħ

72

Human; enzyme; acetyl-Coenzyme A-carboxylase-alpha; ACC-alpha; cancer; breast; ovary; PCR; primer; ss.

Homo sapiens.

WO2002100896-A2

19-DEC-2002.

12-JUN-2002; 2002WO-FR002015

13-JUN-2001; 05-MAR-2002; 2001FR-00007740. 2002FR-00002788.

(CNRS) CNRS CENT

LYON NAT RECH SCI. 1 BERNARD CLAUDE.

Dalla Venezia NL, Magnard CM, Lenoir GM, Sinilnikova-Erard

WPI; 2003-175165/17.

In vitro diagnosis of cancer, particularly break susceptibility, comprises detecting alterations carboxylase alpha gene or protein expression. breast and ovarian tions in the acetyl cancer, or coenzyme A-

Example 1; Page 11; 56pp;

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RESULT 497
AAL60008/c
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Matches 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to human acetyl-Coenzyme A-carboxylase-alpha (ACC-alpha; see ABZ79442), which can be used for in vitro diagnosis of cancer (or of an increased risk of developing it), by detecting ACC-alpha gene mutations or polymorphisms, or altered ACC-alpha protein expression, relative to a control population. The method is particularly used to diagnose cancer, especially of breast or ovary, or for assessing the risk of developing such cancers. The present sequence is a PCR primer, which was used in an example from the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human GH-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAL60008 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20
                                                                                                               The invention relates to growth hormone 1 (GH-1) gene including single nucleotide polymorphisms (SNP). The GH-1 diagnostic polymucleotide is useful as markers for the analysis of a disease, of susceptibility to drug treatment for GH-1 dysfunction or other diseases, or may be included in any complete or partial genetic map of the human genome. GH-1 mutant polypeptides are useful as antagonists of GH-1 hormone action. Polymucleotides encoding these polypeptides are useful in gene therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               27-AUG-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo
                                                                                                                                                                                                                                                           New growth hormone 1 (GH-1) diagnostic polynucleotide, useful as markers for the analysis of a disease, or of susceptibility to drug treatment fo
                                                                                                                                                                                                                                                                                                                                                                              09-NOV-2001; 2001US-0347448P
                                                                                                                                                                                                                                                                                                                                                                                                        07-NOV-2002; 2002WO-US035719
                                                                                                                                                                                                                                                                                                                                                                                                                                  22-MAY-2003
                                                                            Sequence
                                                                                                                                                                                                                       Example 2;
                                                                                                                                                                                                                                                                                                                              Wood
                                                                                                    present
                                                                                                                                                                                                                                                                                                                             LS,
                                                                                                                                                                                                                                              the analysis of a disease, or o dysfunction or other diseases.
                                                                                                                                                                                                                                                                                                   2003-449555/42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            therapy;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          growth
                                                                                                                                                                                                                                                                                                                                                     PHARMACIA & UPJOHN CO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                    Similarity
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                                                                            20
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  CCAAAGTGCTGGGATTACAG
                                                                                                                                                                                                                                                                                                                           Wagner S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    gene
                                                                                                                                                                                                                       Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP; 5 A; 5 C; 5 G; 5 T; 0 U; 0 Other;
                                                                            BP; 4 A; 6 C; 4 G; 6 T; 0 U; 0 Other;
                                                                                                    sequence is a PCR primer used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          hormone 1; GH-1;
                                                                                                                                                                                                                       30;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    amplifying
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              primer; ss
                                                                                                                                                                                                                   74pp;
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                                      2.0%;
                                                                                                                                                                                                                                                                                                                              Parodi LA;
                                                                                                                                                                                                                       English.
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Pred.
                         0;
                                      Score 20;
Pred. No.
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  406
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          single nucleotide polymorphism; SNP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     primer,
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                           Mismatches
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1.1e+03;
                                                   DB 1;
                                       1.1e+03
                                                                                                    for amplifying
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                                                 Length 20;
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                           Indels
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                         ٥,
                                                                                                    therapy.
1 GH-1 gen
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                           Gaps
                                                                                                                                            mutant
                                                                                                                                                       included
                                                                                                      gene
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RESULTY 498
ADD21702/c
ID ADD217
XX
AC ADD217
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AC ADD217
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DT 15-JAN
DX Human
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Human
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Human
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BXAXAXI
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Best Local S
                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention comprises antisense oligonucleotides which are targeted to the human mdm2 gene. The antisense oligonucleotides of the invention are useful for reducing hyperproliferation of human cells. The antisense oligonucleotides are also useful for treating: hyperproliferative disorders (e.g. cancer), psoriasis, fibrosis, atherosclerosis, or restenosis. The antisense oligonucleotides are also useful for modulating apoptosis, and for increasing expression of p21. The present DNA sequence represents a human mdm2 gene antisense oligonucleotide of the invention. The present sequence contains 2'-methoxyethoxy-residues and has a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADD21702
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel antisense compound targeted to 5' untranslated region, region, or intron:exon junction of nucleic acid molecule encuseful for treating e.g. cancer, psoriasis or restenosis by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human mdm2 antisense oligonucleotide #265.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADD21702;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Miraglia LJ,
Manoharan M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO2003048315-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 4 A; 7 C; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 9; SEQ ID NO 267; 289pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2003-577263/54.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  04-DEC-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       02-DEC-2002; 2002WO-US038281
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2'-methoxyethoxy-residue; phosphorothioate backbone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   hyperproliferative disorder; cancer; psoriasis; fil atherosclerosis; restenosis; apoptosis modulation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         antisense
                                                  15-JAN-2004
                                                                                                     ADD21701;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              phosphorothioate backbone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (ISIS-) ISIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12-JUN-2003
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                                                                                                                                                        ADD21701
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                                                                                                                                                                                                                                                                                                                 388 CAAAGTGCTGGGATTACAGG 407
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20
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                                                                                                                                                                                                                                                                                                                                                                                       20;
                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CCAAAGTGCTGGGATTACAG
                                                                                                                                                        standard;
                                                                                                                                                                                                                                                                                    CAAAGTGCTGGGATTACAGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         oligonucleotide; human; mdm2; hyperproliferation;
                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PHARM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2001US-00005344.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first
                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nero PS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INC
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                                                                                                                                                        DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               entry)
                                                                                                                                                                                                                                                                                                                                                                                                               2.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           G; 6 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                               Score 20;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                  .1e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Koller
                                                                                                                                                                                                                                                                                                                                                                                             Indels
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encoding mdm2,
by inhibiting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Chiang
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                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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Human mdm2

antisense oligonucleotide #264.

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RESULT 500
ADD21677/c
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention comprises antisense oligonucleotides which are targeted to the human mdm2 gene. The antisense oligonucleotides of the invention are useful for reducing hyperproliferation of human cells. The antisense oligonucleotides are also useful for treating: hyperproliferative disorders (e.g. cancer), psoriasis, fibrosis, atherosclerosis, or restenosis. The antisense oligonucleotides are also useful for modulating apoptosis, and for increasing expression of p21. The present DNA sequence represents a human mdm2 gene antisense oligonucleotide of the invention. The present sequence contains 2'-methoxyethoxy-residues and has a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo
                                                                                                        antisense oligonucleotide; human; mdm2; hyperproliferation; hyperproliferative disorder; cancer; psoriasis; fibrosis; atherosclerosis; restenosis; apoptosis modulation; p21; ss;
                                                                                                                                                               Human mdm2 antisense oligonucleotide #240.
                                                                                                                                                                                                                                                     ADD21677
                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 9; SEQ ID NO 266; 289pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel antisense compound targeted to 5' untranslated region, coding region, or intron:exon junction of nucleic acid molecule encoding mdm2, useful for treating e.g. cancer, psoriasis or restenosis by inhibiting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2003-577263/54.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Miraglia LJ,
Manoharan M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                04-DEC-2001; 2001US-00005344
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      antisense oligonucleotide; human; mdm2; hyperprolife hyperproliferative disorder; cancer; psoriasis; fibr atherosclerrosis; restenosis; apoptosis modulation; p2'.-methoxyethoxy-residue; phosphorothioate backbone.
                                        WO2003048315-A2
                                                                                             2'-methoxyethoxy-residue; phosphorothioate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            02-DEC-2002; 2002WO-US038281
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         12-JUN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                              phosphorothioate backbone.
                                                                                                                                                                                                                                                                                                                                                  851
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                                                                                                                                                                                                                                                                                                                         20
                                                                                                                                                                                                                                                                                                                                                                              20;
                                                                                                                                                                                                                                                    standard;
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                                                                                                                                                                                                                                                                                                                                                                           2.0%; Solitarity 100.0%; If Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                  BP; 3 A; 7 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                            (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nero PS,
                                                                                                                                                                                                                                                    DNA;
                                                                                                                                                                                            entry)
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0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                           1.1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                       DB 1;
                                                                                                                                                                                                                                                                                                                                                                              0
                                                                                                                                                                                                                                                                                                                                                                                                       Length 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     fibrosis;
                                                                                                                                                                                                                                                                                                                                                                              Indels
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RESULT 501
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ID ABZ979
XX ABZ979
XX ABZ979
XX ABZ979
XX Human
XX Homo s
XX Ing 1
XX ADZ002
XX Homo s
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                 Nyce JW,
Miller S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         the human mdm2 gene. The antisense oligonucleotides of the invention are useful for reducing hyperproliferation of human cells. The antisense oligonucleotides are also useful for treating: hyperproliferative disorders (e.g. cancer), psoriasis, fibrosis, atherosclerosis, or restenosis. The antisense oligonucleotides are also useful for modulating apoptosis, and for increasing expression of p21. The present DNA sequence represents a human mdm2 gene antisense oligonucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel antisense compound targeted to 5' untranslated region, region, or intron: exon junction of nucleic acid molecule encuseful for treating e.g. cancer, psoriasis or restenosis by i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human RANTES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present sequence contains phosphorothioate backbone.
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Manoharan M;
                                                                                                                                                                            24-APR-2001; 2001US-0286137P
                                                                                                                                                                                                                             23-APR-2002;
                                                                                                                                                                                                                                                                                  31-OCT-2002.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              02-DEC-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local
                                                                                                                                                                                                                                                                                                                                                                                     sapiens.
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20; Conserv
                                                                                                                             EPIGENESIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  standard; DNA;
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                                                 Li Y,
Tang
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2.0%;
llarity 100.0%;
Conservative
                                                                                                                                                                                                                                2002WO-US013135
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           oligonucleotide sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
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                                                 Sandrasagra A,
L, Shahabuddin
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                                                                                                                                                                                                                                                                                                                                                                                                                                    respiratory
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           7 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       242; 289pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0,
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            956
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                                                                           Katz
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ₽Þ,
                                                                           Pabalan
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by inhibiting
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WPI; 2003-229219/22

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or

Disclosure; SEQ ID NO 13153; 872pp; English

The invention relates to a novel pharmaceutical composition, which has first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of

ubiquinone

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CC immunosuppressive, and cytostatic activity. The composition may have a continuing and/or cc antiinflammatory steroid and ubiquinone. A composition of the invention cc antiinflammatory steroid and ubiquinone. A composition of the invention cc antiinflammatory, antiallergic, antiasthmatic, hypotensive, cc immunosuppressive, and cytostatic activity. The composition may have a cc use in antisense gene therapy. The composition is useful for treating or cc preventing a respiratory, lung or malignant disease or condition, also cc for enhancing the prophylactic or therapeutic respiratory effect of an cantiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or clung surfactant in a subject tissue, or treating bronchoconstriction, contein the sequence data for this patent is not represented in the printed case of the composition, but was obtained in electronic format directly from WIPO case for wine information, which is held not remembered.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABZ99076
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 502
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Best Local S
Matches 20
                                                                                                                                                                                                                                                                                    Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                  Human PDE4C oligonucleotide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                  17-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABZ99076 standard; DNA; 20
                                                                                                                                                                         31-OCT-2002.
                                                                                                                                                                                                         WO200285308-A2
                                                                                                                                                                                                                                     Homo sapiens
                                              Nyce JW,
                                                                                                                                           23-APR-2002; 2002WO-US013135
                                                                                                          24-APR-2001; 2001US-0286137P
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                                                                              EPIGENESIS PHARM INC
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                          Li Y, Sa
, Tang L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP; 4 A;
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                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                               Sandrasagra A,
L, Shahabuddin
                                                                                                                                                                                                                                                                     respiratory disease; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 20;
Pred. No.
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                                                Katz E,
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                                                Pabalan J,
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Crimitiation relates to a novel pharmaceutical composition, which has a crive agent comprising an oligonucleotide antisense to the crimitiation codon, coding region, 5' or 3' end genomic flanking regions, coding region, 5' or 3' end genomic flanking regions, coding region, 5' or 3' end genomic flanking regions, coding and 3' intron-exon junctions, or regions within 2-10 nucleotides of codinctions of genes encoding a polypeptide associated with lung and/or coding antiangual active agent comprising an companient antisense and ubiquinone. A composition of the invention code in antisense gene therapy. The composition is useful for treating or greventing a respiratory, lung or malignant disease or condition, also composition genetication, also containflammatory steroid in a subject, for reducing or depleting levels of an antiinflammatory steroid in a subject, for reducing or depleting levels of or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject, s tissue, or treating bronchoconstriction, composition in the sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at fig. at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pharmaceutical composition for treating ailments associated with impaired Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; SEQ ID NO 14318; 872pp; English
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Matches Query Match Local 379 TCAGCCTCCCAAAGTGCTGG 398 20; Similarity Conservative 2.0%; 0, Score 20; Pred. No. Mismatches 1.1e+03; DB 1; 0 Length 20; 0 Gaps

0

Sequence 20 BP; 4 A; 7 C; 5 G; 4 T; 0 U; 0 Other;

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ABZ98014 RESULT 503 Human 17-OCT-2003 ABZ98014 standard; DNA; 20 BP RANTES oligonucleotide sequence. (first entry)

Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; Nyce JW, Miller S, Homo sapiens. WPI; 2003-229219/22 24-APR-2001; 2001US-0286137P 23-APR-2002; 2002WO-US013135 Pung (EPIG-) EPIGENESIS PHARM WO200285308-A2 inflammation; respiratory Li Y, Sa Tang L, Sandrasagra A, Shahabuddin s Katz μ Pabalan Ġ Aguilar

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Cc Note: The sequence data for this patent is not represented in the printed case for wine int/cnih/subject and core sequences.
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                                 Nyce JW,
Miller S,
                                                                                                                         24-APR-2001;
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                                                                                                                                                                                                                                                                                                                     antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antiesthe gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
                                                                                                                                                                                                                                                                                                                                                                                                                               Human PDE4C oligonucleotide sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its respiration, has oligo(s) antisense to specific gene(s) or its responding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                   Homo sapiens
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                               Li Y,
Tang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     n codon, coding region, 5' or 3' end genomic flanking regions, intron-exon junctions, or regions within 2-10 nucleotides of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                       2001US-0286137P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
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                                 Sandrasagra A,
, Shahabuddin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        t comprising an oligonucleotide antisense to the coding region, 5' or 3' end genomic flat.
                                                                                                                                                                                                                                                                                                      respiratory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%;
                                                                                                                                                                                                                                                                                                                                                                                            dysfunction; nasal airway dysfunction;
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Pred. No.
                                                                                                                                                                                                                                                                                                      disease; ds.
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has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its responding RNAs, and glucocorticoid or non-glucocorticoid steroid or continuity.
                                                                                                                                                                                                                                                                                                                                                                                              nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                            junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking region, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQ ID NO 14297; 872pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              relates to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           a novel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           pharmaceutical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              composition,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              which has a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       regions,
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Query Match Sequence Similarity 20 BP; 3 A; 5 C; 9 G; 3 T; 0 U; 0 Other; 2.0%; Score 20; Pred. No. DB 1; Length 20;

ftp.wipo.int/pub/published_pct_sequences

멼 Ś 643 CCCAGGCTGGAGTGCAGTGG 662 20

Matches

20;

Conservative

, ,

Mismatches

Indels

0

Gaps

0

1.1e+03;

Local

RESULT 505 ABZ99075 ABZ99075 standard; DNA; 20

Human PDE4C oligonucleotide sequence.

17-OCT-2003

(first entry)

antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene thera Human; antisense; lung dysfunction; nasal airway dysfunction;

Homo sapiens

inflammation;

respiratory disease; ds.

31-OCT-2002

23-APR-2002; 2002WO-US013135

24-APR-2001; 2001US-0286137P

EPIGENESIS PHARM INC.

Nyce JW, Miller S, Li Y, Tang ŗ Sandrasagra A, L, Shahabuddin Katz S; μ Pabalan Ç Aguilar

WPI; 2003-229219/22

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or

The invention relates to a novel pharmaceutical composition, which has first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of

Disclosure; SEQ ID NO 14317; 872pp; English

ubiquinone

cc junctions of genes encoding a polypeptide associated with lung and/or cc nasal airway dysfunction and a second active agent comprising an cc antiinflammatory steroid and ubiquinone. A composition of the invention cc immunosuppressive, antiallergic, antiasthmatic, hypotensive, acc immunosuppressive, and cytostatic activity. The composition may have a cc use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also cf or enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing levels of of or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, containing inflammation, lung allergies, or a respiratory disease or condition.

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RESULT 506
ABZ92715
ID ABZ927
XX ABZ927
XX Human
DT 17-OCT
XX Human
XX Human
KW Antiir
KW Antiir
KW Antiir
KW Antiir
KW Adenoo
KW Lung :
XX Homo
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                                              Nyce JW,
Miller S,
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                                                                                                                  EPIGENESIS PHARM INC
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                                            Li Y, Sa
Tang L,
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                                                 Sandrasagra A,
L, Shahabuddin
                                                   S
                                                                        Katz E,
                                                                        Pabalan J,
                                                                        Aguilar
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Nyce JW, Miller S,

Li Y, Tang

Sandrasagra A, L, Shahabuddin

S

Katz ŭ

Pabalan

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Aguilar D;

(EPIG-)

EPIGENESIS PHARM INC

24-APR-2001; 2001US-0286137P

WPI; 2003-229219/22

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RESULT 507
ABZ92716
ID ABZ927
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CC initiation coding region, 5 or 3 end genomic flanking regions, CC initiation coding, coding region, 5 or 3 end genomic flanking regions, CC initiation coding, coding regions within 2-10 nucleotides of CC junctions of genes encoding a polypeptide associated with lung and/or CC nasal airway dysfunction and a second active agent comprising an CC antiinflammatory steroid and ubiquinone. A composition of the invention CC inmunosuppressive, antiallergic, antiasthmatic, hypotensive, CC immunosuppressive, and cytostatic activity. The composition may have a CC use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also CC for enhancing the prophylactic or therapeutic respiratory effect of an CC antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine CC lung inflammation, lung allergies, or a respiratory disease or condition. CC lung surfactant in a subject's tissue, or treating bronchoconstriction, CC lung inflammation, lung allergies, or a respiratory disease or condition. CC specification, but was obtained in electronic format directly from WIPO can be seen the condition of the printed of the prin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; antisense; lung dysfunction; nasal airway dysfunction; antianflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
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Matches Query Match

Similarity

2.0%; Score 20; DB 100.0%; Pred. No. 1. ative 0; Mismatches

DB 1; 1.1e+03;

Length 20; Indels

0;

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Conservative

Sequence 20

BP; 2 A; 12 C; 2 G; 4 T; 0 U; 0 Other;

lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences

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                             Nyce JW,
Miller S,
WPI; 2003-229219/22
                                                                                                             24-APR-2001; 2001US-0286137P
                                                                                                                                              23-APR-2002;
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                                                                                                                                                                                                                                                                                            antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
                                                                                                                                                                                                                                                                                                                                            Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory; antiallergic;
                                                                                                                                                                                                                                                                                                                                                                                            Human PDE4C oligonucleotide sequence.
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Tang
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                             Sandrasagra A,
J, Shahabuddin
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has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                   junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention antiinflammatory steroid and ubiquinone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to a novel pharmaceutical composition, which I first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking region, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
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lung and/or
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S Matches Query Match Local 772 TTGTATTTTTAGTAGAGATG 791 20; Similarity Conservative 2.0%; 0; Score 20; Pred. No. Mismatches B .1e+03; 1; Length Indels 0 Gaps 0

Sequence

ftp.wipo.int/pub/published_pct_sequences

20 BP; 5 A; 0 C; 5 G; 10 T; 0

U; 0 Other;

RESULT 509 ABX94882 standard; cDNA; 20

Human MBHBIK receptor P2Y34 PCR primer

(first entry)

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TTGTATTTTAGTAGAGATG

20

G protein-coupled receptor; immunomodulatory; gastrointestinal; antiinflammatory; cardiovascular; gene therapy; intestinal function; blood pressure; blood flow; blood coagulation; haematopoiesis; interleukin; prostaglandin; inflammation; neuronal function; cell growth; DE10142478-A1 Homo sapiens. differentiation; Human; receptor; MBHBIK receptor; P2Y34 receptor; chromosome primer; ss.

20-MAR-2003.

31-AUG-2001; 2001DE-01042478

31-AUG-2001; 2001DE-01042478

(BRUE/) BRUESS M. BOENISCH Ξ

VON

KUEGELGEN

Bruess M, Boenisch H, Von Kuegelgen I;

2003-383212/37

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                the invention has immunomobilatory, gastrointestinal, antiinflammatory CC and cardiovascular activity and can be used for gene therapy. The creeptor described in the disclosure may be implicated in regulation of CC intestinal function, blood pressure, blood flow through organs and CC regions of the body; blood coagulation, haematopoiesis and immune CC reactions; release of interleukins and prostaglandins, i.e. in CC inflammation; modulation of neuronal function and cell growth and CC differentiation. The polynucleotide of the invention which encodes a GC grotein-coupled receptor, and also its related cDNA, mRNA, protein, CC antibodies and oligonucleotides, are useful in the diagnosis and CC treatment of diseases associated with abnormal levels of P2Y34 expression CC; in screening assays for modulators, potentifing diseases associated CC with abnormal expression of P2Y34. This sequence represents a PCR primer CC used to amplify the gene encoding the human P2Y34 protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 510
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Best Local S
Matches 20
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   Novel isolated or purified polypeptide encoded by genes associated with intestinal epithelium or M cell development, differentiation or function useful for treating autoimmune diseases and infectious diseases.
                                                                                                                                                                                                                                                                                            04-APR-2001; 2001US-0281416P
                                                                                                                                                                                                                                                                                                                                                       04-APR-2002; 2002WO-US010873
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                incompatibility; ss; human; PCR; primer.
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                The invention describes a method of determining the ethnic origin of a male comprising obtaining a nucleic acid sample from the male, and identifying at least two polymorphic markers in the nucleic acid sample indicative of the ethnic origin of the male, using at least one primer pair from the primer pairs given in the specification. Also described is a method of: identifying polymorphic sites in a nucleic acid; a kit for determining the ethnic origin of an individual; determining the ethnic origin of an individual; determining the sethnic origin of a human male individual; an isolated nucleic acid segment of a primer of a nucleic acid segment of a primer origin of a human male individual; an isolated nucleic acid segment of a primer origin of a human male individual; an isolated nucleic acid segment of a primer origin of a human male individual; an isolated nucleic acid segment of a primer origin of a human male individual; an isolated nucleic acid segment of a primer origin of a human male individual; an isolated nucleic acid segment of a primer origin of a human male individual; an isolated nucleic acid segment of a primer origin of a human male individual; an isolated nucleic acid; a primer origin of a primer origin of a human male individual; an isolated nucleic acid; a primer origin of a primer origin of a primer or origin or or
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                                                                                                                                                                                                                                                                                                  Determining the ethnic origin of a male by obtaining a nucleic acid sample from the male and identifying at least two polymorphic market the nucleic acid sample indicative of the ethnic origin of the male
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               r cnromosome; paternity non-recombining region;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ethnic origin determination; polymorphic site determination; Y chromosome; paternity testing; forensic; diagnosis; non-recombining region; human; NRV; polymorphic fragment; ds
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                                                                                                                                                                                                                                                   Page 65; 74pp; English
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CC The invention describes a method of determining the ethnic origin of a CC male comprising obtaining a nucleic acid sample from the male, and CC identifying at least two polymorphic markers in the nucleic acid sample CC indicative of the ethnic origin of the male, using at least one primer CC pair from the primer pairs given in the specification. Also described is CC a method of: identifying polymorphic sites in a nucleic acid; a kit for CC determining the ethnic origin of an individual; determining the ethnic cridin of a human male individual; an isolated nucleic acid segment of a human y chromosome comprising at least 10 contiguous bases including at CC least one of the polymorphic sites given in the specification; nucleic acid primer pairs for amplifying polymorphic regions of the y chromosome CC individual. The method is useful for determining the ethnic origin of a complete compart of the paternity of a human male individual. The method is useful for determining the ethnic origin of a complete compart of the paternity testing, for forensic studies or for diagnosis. This sequence represents a fragment of the non-recombining region of the human cc y chromosome (NRY) comprising a polymorphism that can be used to
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Query Match
Best Local Similarity
Matches 20; Conserv

Conservative

0,

Mismatches

100.0%; 2.0%;

Score 20; Pred. No.

DB 1;

Length 20; Indels

1.1e+03

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Gaps

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RESULT 513
ADM65575/c
ID ADM655
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                        The invention describes a method of determining the ethnic origin of a CC male comprising obtaining a nucleic acid sample from the male, and CC identifying at least two polymorphic markers in the nucleic acid sample CC identifying at least two polymorphic markers in the nucleic acid sample CC indicative of the ethnic origin of the male, using at least one primer CC as method of: identifying polymorphic sites in a nucleic acid; a kit for CC determining the ethnic origin of an individual; determining the ethnic origin of an isolated nucleic acid segment of a fundam Y chromosome comprising at least 10 contiguous bases including at CC least one of the polymorphic sites given in the specification; nucleic acid primer pairs for amplifying polymorphic regions of the Y chromosome CC individual. The method is useful for determining the ethnic origin of a CC male, for paternity testing, for forensic studies or for diagnosis. This sequence represents a primer used to detect polymorphisms in the non-recombining region of the human Y chromosome (NRY).
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ethnic origin determination; polymorphic site determination; Y chromosome; paternity testing; forensic; diagnosis; non-recombining region; human; NRY; PCR; primer; ss.
                                                                                                                                                                                                                                                                                 Claim
                                                                                                                                                                                                                                                                                                           Determining the ethnic origin of a male by obtaining a nucleic acid sample from the male and identifying at least two polymorphic markers in the nucleic acid sample indicative of the ethnic origin of the male.
                                                                                                                                                                                                                                                                                                                                                                                                 Oefner PJ,
                                                                                                                                                                                                                                                                                                                                                                                                                           (OEFN/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-NOV-2000; 2000US-0245355P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-NOV-2001; 2001US-00002623.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NRY polymorphism detection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            03-JUN-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       determine ethnic origin of a male.
                                                                                                                                                                                                                                                                                                                                                                      WPI; 2003-843259/78.
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                                                                                                                                                                                                                                                                              24; Page 54; 74pp; English.
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l Similarity 100.0%;
20; Conservative
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UNDERHILL P
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B₽;
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   7
C; 3
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 0 Other;
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RESULT 514
ADM65745
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                                                                                                                                                                                                                    RESULT 515
ADM65578/c
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                                                                                                                                                                                                                                                                    The invention describes a method of determining the ethnic origin of a male comprising obtaining a nucleic acid sample from the male, and conditiving at least two polymorphic markers in the nucleic acid sample indicative of the ethnic origin of the male, using at least one primer condition of the male, using at least one primer condition of the specification. Also described is a method of: identifying polymorphic sites in a nucleic acid; a kit for contigin of a human male individual; an individual; determining the ethnic origin of an individual; determining the ethnic corigin of a human with the polymorphic sites given in the specification; nucleic acid grimer pairs for amplifying polymorphic regions of the y chromosome comprising at least 10 contiguous bases including at least one of the polymorphic sites given in the specification; nucleic carid primer pairs for amplifying polymorphic regions of the y chromosome carid in the specification; and determining the paternity of a human male individual. The method is useful for determining the ethnic origin of a male, for paternity testing, for forensic studies or for diagnosis. This sequence represents a fragment of the non-recombining region of the human cycle y chromosome (NRY) comprising a polymorphism that can be used to
                                                                                                                                              Matches
                                                                                                                                                               Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Determining the ethnic origin of a male by obtaining a nucleic acid sample from the male and identifying at least two polymorphic markers the nucleic acid sample indicative of the ethnic origin of the male.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Y chromosome; paternity testing; forensic; diagnosis; non-recombining region; human; NRY; polymorphic fragment; ds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ethnic origin determination; polymorphic site determination;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human Y chromosome non-recombining region polymorphic fragment
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                                                                                                                                                                                                                  Sequence 20 BP; 6 A;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (UNDE/) UNDERHILL P A.
                                                                                                                                                                                                                                                      chromosome (NRY) comprising a polymorphism that termine ethnic origin of a male.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              389
                                                                                                          388 CAAAGTGCTGGGATTACAGG 407
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                                                                                                                                                               Similarity
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                                                                                                                                                               2.0%;
                                                                                                                                                                                                                    3 C; 7 G; 4 T; 0
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                                                                                                                                              0;
                                                                                                                                                               Score 20; DB 1; L
Pred. No. 1.1e+03;
                                                                          20
                                                                                                                                                Mismatches
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RESULT 516
ADM34330/c
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Best Local S
Matches 20
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                  Human cryopyrin cDNA sequence primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 24; Page 55; 74pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2003-843259/78.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-NOV-2000; 2000US-0245355P
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                                                                 03-JUN-2004
                                                                                                             ADM34330;
                                                                                                                                                          ADM34330 standard; DNA; 20
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                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
                                                                                                                                                                                                                                                                                                             389 AAAGTGCTGGGATTACAGGC 408
                                                                                                                                                                                                                                                                     20
                                                                                                                                                                                                                                                                                                                                                            20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP;
                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                   (first entry)
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                                                                                                                                                                                                                                                                                                                                                     100.0%; --
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                                                                                                                                                                                                                                                                                                                                                                                                        2.0%;
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                         #2
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RESULT 517
ABD32099
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Best Local S
Matches 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         acid sequence of wild type cryopyrin of 1034 amino acids, with the proviso that amino acid 198 is not Val, amino acid 352 is not Ala, amino acid 434 is not Ala, amino acid 627 is not Glu, or amino acid 703 is not Gln. The methods are useful for determining the presence of a disorder, treating inflammation, Familial cold urticaria/familial cold autoinflammatory syndrome (FCU/FCAS) or Muckle Wells Syndrome (MWS), and identifying a substance useful in modulating binding of a cryopyrin protein to a second protein. The oligonucleotide is useful in diagnosing a disorder characterized by an aberrant CIASI gene. This sequence corresponds to a primer used to amplify the cryopyrin gene of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New isolated cryopyrin protein and encoding nucleic acid, useful for diagnosing and treating inflammatory disorders, in particular familial cold autoinflammatory syndrome and/or Muckle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hoffman H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        05-OCT-2001; 2001US-0327728P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         04-OCT-2002; 2002WO-US031502
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ss; primer; antiinflammatory; cryopyrin; inflammation;
Familial cold urticaria; familial cold autoinflammatory syndrome;
                                                                                                                                                            Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer;
                                                                                                                                                                                                              Human PDE4C-derived oligonucleotide SEQ ID 14310.
                                                                                                                                                                                                                                                                              ABD32099
                                                                                                                                                                                                                                                                                                               ABD32099 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20
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                                                             respiratory distress syndrome; allergic rhinitis; pulmonary emphysema; chronic obstructive pulmonary disease; cancer; br pulmonary transplantation rejection; ss; primer.
                                                                                                         surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vascoonstriction;
                                                                                                                                                                                                                                              29-JUL-2004
                              Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       invention relates to a novel isolated protein (I) comprises the amino sequence of wild type cryopyrin of 1034 amino acids. with the
                                                                                                                                                                                                                                                                                                                                                                                                                              667
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                                                                                                                                                                                                                                                                                                               DNA;
                                                                                                                                                                                                                                              entry)
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Pred. No.
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                                                                                            hypertension;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            0
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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WO200285309-A2

RESULT 518 ABD31045

ABD31045 standard; DNA; 20

吊 8

Matches Query Match Best Local

20;

Conservative

0;

Mismatches

0

Gaps

0

Similarity

2.0%;

Score 20; DB 1; Lo Pred. No. 1.1e+03;

Length 20;

772 ш

TTGTATTTTTAGTAGAGATG 791 TTGTATTTTAGTAGAGATG 20

SEXEXEX

29-JUL-2004 (first entry)

Human RANTES-derived oligonucleotide

SEQ

ID 13256

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cc surfactant depletion or hyposecretion, when administered to a mammal. The capture of coligonucleotides are derived from a gene encoding or regulating conjugation of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery constructions for adding a carrier and for use of the invention has antiallergic, antiinflammatory, antiasthmatic, considered in the invention has antiallergic, antiinflammatory, antiasthmatic, considered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The conformation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, considered inflammation, allergies, asthma, impeded respiration, respiratory hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to the oligonucleotides into products that free adenosine into the system ceg., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, all reducing adenosine sensitivity, levels of adenosine (A)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and
Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This invention describes a novel composition (a) a first active
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 15; SEQ ID NO 14310;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         bronchodilating agent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    24-APR-2001; 2001US-0286036P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          31-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (EPIG-) EPIGENESIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Li 1,
Tang L,
20
BP; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2002WO-US013143
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, Shahabuddin
A; 0 C; 5 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             763pp; English
10 T; 0 U;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Katz E,
0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         allergies
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      receptors,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        agent,
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respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distrass syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                   Human; antisense;
                                                                                                                                                                                                                                                                                                                                                                                                                   bronchoconstriction; allergy; hyposecretion; pain;
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sapiens.

WO200285309-A2

23-APR-2002; 2002WO-US013143

24-APR-2001; 2001US-0286036P

(EPIG-) EPIGENESIS PHARM INC

S Li Y, Tang Ļ Sandrasagra A, Shahabuddin ŝ Katz Ħ Pabalan Ç Aguilar D;

WPI; 2003-093058/08

Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted bronchodilating agent. nucleic acids associated with lung airway or lung dysfunction, ç

SEQ ID NO 13256; 763pp; English

CC comprising oligonucleotides, effective for alleviating
CC pronchoconstriction, respiratory tract inflammation, allergies and
CC reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors,
CC surfactant depletion or hyposecretion, when administered to a mammal. The
CC oligonucleotides are derived from a gene encoding or regulating
CC expression of a target polypeptide associated with lung airway or lung
CC dysfunction or cancer and can be anti-sense to the corresponding mRNA.
CC The invention also describes a kit, that comprises: (a) a delivery
CC instructions for adding a carrier and for use of the kit. The composition
CC of the invention has antiallergic, antiinflammatory, antiasthmatic,
CC analgesic, hypotensive, immunosuppressive and cytostatic activity, is a
CC beta-adrenergic agonist. The composition is useful for preventing or
CC treating a respiratory, lung or malignant disease. The administered
CC composition comprises oligo and is administered to reduce the production
CC or availability, or to increase the degradation of the target mRNA or to
CC reduce the amount of target polypeptide present in the lungs. The
CC inflammation, allergies and/or burnchoconstriction and/or lung
CC with a disease or condition such as pulmonary vasoconstriction
CC inflammation, allergies, asthma, impeded respiration, respiratory
CC transplantation rejection, pulmonary infections, bronchitis or cancer.
CC The reduced adenosine content of the anti-sense oligos corresponding to
CC the oligonary into the target RNA serves to prevent the breakdown of
CC the oligonary into the administic the breakdown of
CC the oligonary into the administ the forms administering the corresponding to
CC the oligonary the content of the anti-sense oligos corresponding to This invention describes oligonucleotides into products that ., lung, brain, heart, kidney, etc, t vent any unwanted effects due to it a novel composition (a) a first active agent, tissue environment free adenosine into the system

20 B₽; ω P 4 C; 10 ç, ω Τ; 0 Ģ; 0 Other;

Query Match Best Local S Matches 20 Similarity 20; Conser Conservative 2.0%; 0 Score 20; Pred. No. Mismatches DB 1; 1.1e+03 Length 20; Indels 0 Gaps

0

밁 S

> RESULT 519 ABD30942 Human RANTES-derived oligonucleotide SEQ ABD30942; ABD30942 standard; (first entry) DNA; 20

respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; systic fibrosis; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; analgesic; hypotensive; immunosuppressive; cytostatic; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; Human; antisense; pulmonary transplantation rejection; ss; primer bronchoconstriction; allergy; pulmonary vasoconstriction; hyposecretion; pain;

ID 13153

Homo sapiens.

WO200285309-A2

31-OCT-2002

2002WO-US013143

24-APR-2001; 2001US-0286036P

(EPIG-) EPIGENESIS PHARM INC

ß Li Y, Tang ŗ Sandrasagra A, C, Shahabuddin Katz E, Pabalan ç Aguilar D;

Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted nucleic acids associated with lung airway or lung dysfunction, and bronchodilating agent. ö

Claim 15; SEQ ID NO 13153; 763pp; English

CC comprising of injunction, respiratory tract inflammation, allergies and composition gadenosine sensitivity, levels of adenosine (A) or (A) receptors, comprising of ignnucleotides, effective for alleviating controlled adenosine sensitivity, levels of adenosine (A) or (A) receptors, consider the depletion or hyposecretion, when administered to a mammal. The coligonucleotides are derived from a gene encoding or regulating controlled at target polypeptide associated with lung airway or lung controlled internation also describes a kit, that comprises: (a) a delivery controlled internation also describes a kit, that comprises: (a) a delivery controlled internation and for use of the invention has antiallergic, antiinflammatory, antiasthmatic, controlled in hypotensive, immunosuppressive and cytostatic activity, is a composition for adding a carrier and for use of the kit. The composition composition are septiatory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vascoonstriction, confidence, palin, cystic fiborosis, allergic ribinitis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary cancer. hypertension, emphysema, GRIGHLE OFFICERS, bronchitis or cancer. transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to the reduced adenosine content the serves to prevent the breakdown of thymidines present in the target RNA serves to prevent the breakdown of thymidines present in the target RNA serves to prevent the breakdown of thymidines present in the system. This invention describes a novel composition (a) a first active agent,

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RESULT 520
ABD32107
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Best Local
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prevent
                                           comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the ollgonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a
                                                                                                                                                                                                                                                                                                                        Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted nucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; antisense;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human PDE4C-derived oligonucleotide SEQ ID 14318.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABD32107,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABD32107 standard;
 beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the product
                                                                                                                                                                                                                                                                         Claim 15; SEQ ID NO 14318; 763pp; English
                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2003-093058/08.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      24-APR-2001; 2001US-0286036P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      23-APR-2002; 2002WO-US013143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      31-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO200285309-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   pulmonary transplantation rejection; ss; primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                29-JUL-2004
                                                                                                                                                                                                                                             This invention describes a novel composition (a) a first active
                                                                                                                                                                                                                                                                                                           bronchodilating agent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (EPIG-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     .g., lung,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EPIGENESIS PHARM INC
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                                                                                                                                                                                                                                                                                                                                                                                                                    Li Y,
Tang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ilarity 100.0%;
Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 4 A; 7 C; 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        unwanted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       brain, heart, kidney, etc, tissue environment
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                                                                                                                                                                                                                                                                                                                                                                                                                      ŗ
                                                                                                                                                                                                                                                                                                                                                                                                                         Sandrasagra A,
L, Shahabuddin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                bronchoconstriction; allergy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        effects due
oligo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        G; 2 T; 0 U;
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                                                                                                                                                                                                                                                                                                                                                                                                                                       Katz E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1.1e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  hyposecretion;
                                                                                                                                                                                                                                                                                                                                                                                                                                        ŗ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     and thereby,
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   production
                                                                                                                                                                                                                                                                                                                            and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  pain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                               agent
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pulmonary obstruction, and/or bronchoconstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonnucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to
Sequence
                                                                                                    prevent any unwanted effects due to it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      reduce the amount
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      or availability,
20 BP; 4 A; 7 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 or to increase the degradation of the target mRNA t of target polypeptide present in the lungs. The
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Query Match
Best Local S
                                 Matches
             379 TCAGCCTCCCAAAGTGCTGG 398
                                l Similarity
20; Conserv
ш
TCAGCCTCCCAAAGTGCTGG
                                 Conservative
                                          2.0%;
                                 0;
                                                   Score 20;
20
                                           Pred. No.
                                  Mismatches
                                           1.1e+03;
                                                    DB 1;
                                                   Length
                                  Indels
                                                    20
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                                  Gaps
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RESULT 521
ABD28946
IID ABD289
XX ABD289
XX ABD289
XX ABD289
XX Y
XX HUMANI
KW Fespii
KW Fespii
KW Pulmori
KW Pulmori
KW Pulmori
XX Y
XX HOMO 6
XX POOC
XX HOMO 6
XX HOMO 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antinflammatory; antisthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       N58473-derived oligonucleotide SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          standard; DNA;
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Homo sapiens.

WO200285309-A2

31-OCT-2002

23-APR-2002; 2002WO-US013143.

2001US-0286036P

(EPIG-) EPIGENESIS PHARM INC.

Nyce JW, L Li Y, Tang L, Sandrasagra A, L, Shahabuddin ŝ Katz M Pabalan ۲, Aguilar

Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and bronchodilating

Ħ ö 763pp; English

comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies reducing adenosine sensitivity, levels of adenosine (A) or (A) This invention describes a novel composition (a) a first active receptors, agent,

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CC The invention also describes a kit, that comprises: (a) a delivery CC device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, CC analgesic, hypotensive, immunosuppressive and cytostatic activity, is a CC beta-adrenergic agonist. The composition is useful for preventing or CC treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to CC reduce the amount of target polypeptide present in the lungs. The CC pulmonary obstruction, and/or burnchoconstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, comprises syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to the oligonacleotides into products that free adenosine into the system c.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 522
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local S
Matches 20
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                                                             Nyce JW,
Miller S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis;
WPI; 2003-093058/08
                                                                                                                                                                                                                                  24-APR-2001; 2001US-0286036P
                                                                                                                                                                                                                                                                                                       23-APR-2002; 2002WO-US013143
                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200285309-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; antisense; bronchoconstriction; allergy; hyposecretion; pain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human PDE4C-derived oligonucleotide SEQ ID 14317.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   29-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABD32106 standard;
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                                                         Li Y, Sa
Tang L,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
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                                                                                                 Sandrasagra A,
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                                                                                                                                                                     PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%;
                                                                    Shahabuddin
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Pred. No.
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                                                                    ß
                                                                                                    Katz E,
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                                                                                                    Pabalan J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 20;
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Conjugate to the street from a general encounty of regulations of a target polypeptide associated with lung altrway or lung conversion of a target polypeptide associated with lung altrway or lung conversion also describes a kit, that comprises: (a) a delivery of device, in separate containers, (b) the oligonucleotides, (c) analysic, in separate containers, (b) the oligonucleotides, (c) of the invention has antiallergic, antiinflammatory, antiasthmatic, of analysic, hypotensive, immunosuppressive and cytostatic activity is a beta-adrenergic agonist. The composition is useful for preventing or composition composition composition of the administered composition composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to creduce the amount of target polypeptide present in the lungs. The cumposition, allergies and/or bronchoconstriction and/or lung pulmonary obstruction, and/or bronchoconstriction and/or lung confiammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, confiammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, confiammation, emphysema, chronic obstructive pulmonary disease, pulmonary confiammation, emphysema, chronic obstructive pulmonary disease, pulmonary the reduced adenosine content of the anti-sense oligos corresponding to the oligonucleotides into products that free adenosine into the system of the oligonucleotides into products that free adenosine into the system of the oligonucleotides and effected fine to it
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) recept surfactant depletion or hyposecretion, when administered to a mammal oligonicleotides are derived from a gene encoding or regulating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 15; SEQ ID NO 14317; 763pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 comprising oligonucleotides, effective for alleviating
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Best Matches Query Match Local 369 TCCACCTGCCTCAGCCTCCC 388 Similarity Conservative 2.0%; 0 Score Score 20; Pred. No. Mismatches DB 1; 1.1e+03; Length 20; Indels

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Gaps

0

Sequence 20

any unwanted effects due to

BP; 2 A; 12 C; 2 G; 4 T; 0 U; 0 Other;

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RESULT 523
ABD32086 standard; DNA;
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respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer. Human; antisense; bronchoconstriction; allergy; hyposecretion; pain;

Human PDE4C-derived oligonucleotide SEQ ID 14297.

(first entry)

WO200285309-A2

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Comprising oligonucleotides, effective for alleviating
C bronchoconstriction, respiratory tract inflammation, allergies and
C reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors,
CC surfactant depletion or hyposecretion, when administered to a mammal. The
C oligonucleotides are derived from a gene encoding or regulating
CC expression of a target polypeptide associated with lung airway or lung
CC dysfunction or cancer and can be anti-sense to the corresponding mRNA.
CC The invention also describes a kit, that comprises: (a) a delivery
CC device, in separate containers, (b) the oligonucleotides, (c)
CC instructions for adding a carrier and for use of the kit. The composition
CC of the invention has antiallergic, antiinflammatory, antiasthmatic,
CC analgesic, hypotensive, immunosuppressive and cytostatic activity, is a
CC treating a respiratory, lung or malignant disease. The administered
CC composition comprises oligo and is administered to reduce the production
CC or availability, or to increase the degradation of the target mRNA or to
CC reduce the amount of target polypeptide present in the lungs. The
CC inflammation, allergies and/or surfactant hypoproduction are associated
CC with a disease or condition such as pulmonary vascoconstriction, respiratory
CC distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary
CC transplantation rejection, pulmonary infections, bronchitis or cancer.
CC the reduced adenosine content of the anti-sense oligos corresponding to
CC thymidines present in the target RNA serves to prevent the breakdown of
CC the oligonucleotides into products that free adenosine into the system
CC products and thereby, etc, tissue environment and thereby, to
CC products and content of the anti-sense oligos corresponding to
CC products and content of the anti-sense oligos corresponding to
CC products and content of the anti-sense oligos corresponding to
CC products and content of the anti-sense oligos corresponding to
CC products and content of the anti-sense oligos 
                                                                                                                                                                                                           RESULT 524
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Human; antisense; bronchoconstriction; allergy; hyposecretion;
                                        N58473-derived oligonucleotide SEQ ID 7957
                                                                                     29-JUL-2004
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                                                                                                                                                                  ABD28945 standard;
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Tang
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                                                                                                                                                                                                                                                                                                                                                        Conservative
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                                                                                 (first entry)
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Shahabuddin
                                                                                                                                                                                                                                                                                                                                                                         100.0%;
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S;
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surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
                                                                                                                                                                                                                                                                                                                                                            respiratory tract inflammation; adenosine sensitivity; lung;
                                                                                                                                                                                                                                                                                                                                                                        cancer;
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WO200285309-A2

31-OCT-2002

23-APR-2002; 2002WO-US013143

24-APR-2001; 2001US-0286036P

EPIGENESIS PHARM INC

Nyce Ji Miller е JW, 't S, Li Y, Sa Tang L, Sandrasagra A, L, Shahabuddin , Katz. μ Pabalan ç Aguilar

bronchodilating agent. nucleic Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to acids associated with lung airway or lung dysfunction,

Claim 15; SEQ ID NO 7957; 763pp; English.

containing the content of the many variables and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the antice to prevent the hypertension of the antice to t oligonuclectides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonuclectides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) recepto surfactant depletion or hyposecretion, when administered to a mammal. the oligonucleotides into products that free adenosine in e.g., lung, brain, heart, kidney, etc, tissue environment thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system This invention describes a novel composition (a) a first active lung, nt any unwanted effects due to it receptors, agent, 6

Sequence 20 BP; 4 Þ 9 Ç 4. ç, 3 T; 0 ς, 0 Other;

S Best Lo Query Match Local 540 l Similarity 20; Conser GCCTCAGCCTCCCAAGTAGC 559 2.0%; ۰, Score Pred. Mismatches . 20; DB 1; 1.1e+03 Length 20; ٥, Gaps 0

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RESULT 525
AD180086/c
ID AD1800
XX US2004
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XX DROBEN
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XX The information of the componion of the componion of the componic of the com
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                                                                                                                                                                                                                                                                                         The invention relates to a novel antisense compound of 8-80 nucleobases CC in length targeted to, and which specifically hybridizes with, a nucleic cald molecule encoding transforming growth factor (TGF)-beta 2, and CC inhibits the expression of TGF-beta 2. The invention further relates to: CC a compound 8-80 nucleobases in length that specifically hybridizes with CC at least an 8-nucleobases portion of an active site on a nucleic acid compound encoding TGF-beta 2; a composition comprising the compound and a CC carrier or diluent; inhibiting the expression of TGF-beta 2 in cells or tissues by contacting the cells or tissues with the compound so that CC expression of TGF-beta 2 is inhibited; treating an animal having a CC disease or condition associated with TGF-beta 2 by administering to the compound a therapeutic or prophylactic amount of the compound so that CC expression of TGF-beta 2 is inhibited; and screening an antisense compound. The antisense compound has cytostatic, nootropic, composition and methods are useful for treating a disease or condition associated with TGF-beta 2, such as a hyperproliferative disorder e.g. CC cancer, a neurodegenerative disorder, or a disease or condition involving CC represents an antisense oligonucleotide of the invention.
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                                                                                                                      Matches
                                                                                                                                                      Best Local
                                                                                                                                                                            Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New compounds, particularly antisense oligonucleotides targeted to a nucleic acid encoding TGF-beta 2, useful for treating cancer, a neurodegenerative disorder, or a disease involving hyperactivation of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            antisense; transforming growth factor; TGF; beta 2; TGF-beta 2; cytostatic; nootropic; neuroprotective; immunosuppressive;
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                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (ISIS-) ISIS PHARM INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      hyperproliferative disorder; cancer; neurodegenerative; hyperactivation
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   20
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                                                                                                                                                  Similarity
                                                                                                                                                                                                                                            20
                                                         CGGCCTCCCAAAGTGCTGGG 869
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Freier SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQ ID NO 87; 135pp; English.
                                                                                                                                                                                                                                         BP; 3 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            entry)
                                                                                                                                                  2.0%;
                                                                                                                                                                                                                                         7 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Dobie KW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20
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                                                                                                                      0,
                                                                                                                                                  Score 20;
Pred. No
                                                                                                                         Mismatches
                                                                                                                                                      No.
                                                                                                                                                  DB 1; Le
                                                                                                                            0
                                                                                                                                                                               Length 20;
                                                                                                                            Indels
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                                                                                                                         0,
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                                                                                                                            Gaps
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The invention relates to a novel antisense compound of 8-80 nucleobases CC in length targeted to, and which specifically hybridizes with, a nucleic CC acid molecule encoding transforming growth factor (TGF)-beta 2, and CC inhibits the expression of TGF-beta 2. The invention further relates to: CC a compound 8-80 nucleobases in length that specifically hybridizes with CC at least an 8-nucleobase portion of an active site on a nucleic acid CC molecule encoding TGF-beta 2; a composition comprising the compound and a CC carrier or diluent; inhibiting the expression of TGF-beta 2 in cells or tissues by contacting the cells or tissues with the compound so that CC expression of TGF-beta 2 is inhibited; treating an animal having a CC disease or condition associated with TGF-beta 2 by administering to the CC animal a therapeutic or prophylactic amount of the compound so that CC expression of TGF-beta 2 is inhibited; and screening an antisense compound. The antisense compound has cytostatic, noctropic, composition and methods are useful for treating a disease or condition conservative disorder, or a disease or condition involving cancer, a neurodegenerative disorder, or a disease or condition involving concerts a preferred target DNA region of TGF-beta 2 of the invention.
                                                                                                                     Matches
                                                                                                                                                  Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New compounds, particularly antisense oligonucleotides targeted to nucleic acid encoding TGF-beta 2, useful for treating cancer, a neurodegenerative disorder, or a disease involving hyperactivation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               antisense; transforming growth factor; TGF; beta 2; TGF-beta 2;
cytostatic; nootropic; neuroprotective; immunosuppressive;
hyperproliferative disorder; cancer; neurodegenerative; hyperactivation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human transforming
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 02-JUL-2002; 2002US-00189267.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 16; SEQ ID NO 223; 135pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Monia
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                                                                                                                                                                                                                                           Sequence
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                                                            866 TGGGATTACAGGCGTGAGCC 885
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   \vdash
                                                                                                                     20;
                                                                                                                                                     Similarity
TGGGATTACAGGCGTGAGCC
                                                                                                                                                                                                                                               20
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                                                                                                                         Conservative
                                                                                                                                                                                                                                        BP; 4 A; 4 C; 8 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
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                                                                                                                                                  2.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Dobie KW;
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                                                                                                                         0;
                                                                                                                                                        Score 20;
Pred. No.
                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            English.
                                                                                                                                                        1.1e+03
                                                                                                                                                                                     DB 1;
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                                                                                                                         Indels
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                                                                                                                            <u>0</u>
                                                                                                                            Gaps
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RESULT 527 ADI80087/c ID ADI800 XX AC ADI800

ADI80087 standard; DNA;

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ВP

ADI80087

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RESULT 528
ADI80221
ID ADI802
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AC ADI802
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DT 22-APR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a novel antisense compound of 8-80 nucleobases in length targeted to, and which specifically hybridizes with, a nucleic acid molecule encoding transforming growth factor (TGF) beta 2, and cid inhibits the expression of TGF-beta 2. The invention further relates to: a compound 8-80 nucleobases in length that specifically hybridizes with at least an 8-nucleobase portion of an active site on a nucleoic acid molecule encoding TGF-beta 2; a composition comprising the compound and a carrier or diluent; inhibiting the expression of TGF-beta 2 is inhibited; treating an animal having a CC disease or condition associated with TGF-beta 2 by administering to the expression of TGF-beta 2 is inhibited; treating an animal having a CC disease or condition associated with TGF-beta 2 by administering to the capression of TGF-beta 2 is inhibited; and screening an antisense compound. The antisense compound as cypostatic, nootropic,
                                                                                                                                                                                                                                                 Best Loc
Matches
                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                    Sequence 20
                                                                                                                                                                                                                                                                                                                                                    neuroprotective, and immunosuppressive activities. The compound, composition and methods are useful for treating a disease or condition associated with TGF-beta 2, such as a hyperproliferative disorder e.g. cancer, a neurodegenerative disorder, or a disease or condition involving hyperactivation of an immune response. This polynucleotide sequence represents an antisense oligonucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 15; SEQ ID NO 88; 135pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New compounds, particularly antisense oligonucleotides targeted to a nucleic acid encoding TGF-beta 2, useful for treating cancer, a neurodegenerative disorder, or a disease involving hyperactivation of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2004-081742/08
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                    22-APR-2004
                                                      ADI80221;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     immune; ss; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        antisense; transforming growth factor; TGF; beta 2; TGF-beta 2;
                                                                                     ADI80221 standard;
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                                                                                                                                                                                                                                                                     Local
                                                                                                                                                                                                                966
                                                                                                                                                                               20
                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                TGGGATTACAGGCGTGAGCC 885
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Freier SM,
                                                                                                                                                                                                                                                 ilarity 100.0%;
Conservative
                                                                                                                                                                                                                                                                                                                    BP; 4 A; 8 C; 4 G; 4 T; 0 U; 0 Other;
                    (first entry)
                                                                                       DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         growth factor-beta 2 antisense oligo,
                                                                                                                                                                                                                                                                                  2.0%;
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                                                                                       ₽P
                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                   Score 20;
Pred. No.
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                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                     1.1e+03;
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RESULT 529 ADJ53542

ADJ53542 standard; DNA;

20 ВP 밁 Ś

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850 CGGCCTCCCAAAGTGCTGGG 869

0

CGGCCTCCCAAAGTGCTGGG 20

06-MAY-2004

(first entry

Human; PPP3CB; ss; antisense oligonucleotide; phosphorothioate linkage;

Human PPP3CB DNA antisense oligonucleotide #65.

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acid molecule encoding transforming growth factor (TGF)-beta 2, and CC inhibits the expression of TGF-beta 2. The invention further relates to: CC an compound 8-80 nucleobases in length that specifically hybridizes with CC at least an 8-nucleobase portion of an active site on a nucleic acid compound encoding TGF-beta 2; a composition comprising the compound and a corrier or diluent; inhibiting the expression of TGF-beta 2 in cells or tissues by contacting the cells or tissues with the compound so that compression of TGF-beta 2 is inhibited; treating an animal having a compression of TGF-beta 2 is inhibited; treating an animal having a companial a therapeutic or prophylactic amount of the compound so that compound and attracepture or prophylactic amount of the compound, compound as condition and methods are useful for treating a disease or condition composition and methods are useful for treating a disease or condition associated with TGF-beta 2, such as a hyperproliferative disorder e.g. cascoiated with TGF-beta 3, such as a hyperproliferative disorder e.g. cascoiated with TGF-beta 2, such as a hyperproliferative disorder e.g. cascoiated with TGF-beta 3, such as a hyperproliferative disorder e.g. cascoiated with TGF-beta 2, such as a hyperproliferative disorder e.g. cascoiated with TGF-beta 3, such as a hyperproliferative disorder e.g. cascoiated with TGF-beta 2, such as a hyperproliferative disorder e.g. cascoiated with TGF-beta 3, such as a hyperproliferative disorder e.g. cascoiated with TGF-beta 2, such as a hyperproliferative disorder e.g. cascoiated with TGF-beta 2, such as a hyperproliferative disorder e.g. cascoiated with TGF-beta 2, such as a hyperproliferative disorder e.g. cascoiated with TGF-beta 2, such as a hyperproliferative disorder e.g. cascoiated with TGF-beta 2, such as a hyperproliferative disorder e.g. cascoiated with TGF-beta 2, such as a hyperproliferative disorder e.g. cascoiated with TGF-beta 2, such as a hyperproliferative disorder e.g. cascoiated with TGF-beta 2, such as a hyperprolif
Matches
                                                      Query Match
Best Local
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cytostatic; nootropic; neuroprotective; immunosuppressive;
hyperproliferative disorder; cancer; neurodegenerative; hyperactivation;
                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to a novel in length targeted to, and which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 16; SEQ ID NO 222; 135pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  02-JUL-2002; 2002US-00189267
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         20;
                                                           Similarity
                                                                                                                                                                                                     20 BP; 3
              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PHARM INC
                                                                                                                                                                                                     A; 7 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  growth factor-beta 2 target DNA
                                                      2.0%; Score 20;
100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Dobie KW;
         0;
              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     specifically hybridizes with, a nucleic ing growth factor (TGF)-beta 2, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        antisense compound of 8-80 nucleobases
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                                                                                                            DB 1;
                                                           1.1e+03;
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RESULT 530
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New compounds, particularly antisense oligonucleotides targeted to a nucleic acid encoding PPP3CB, useful for treating an autoimmune disorder, or Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    31-JUL-2002; 2002US-00210723.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2'-O-methoxyethyl sugar moiety; 5-methylcytosine; autoimmune disorder; Alzheimer's disease; immunosuppressive; nootropic; neuroprotective.
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                                                                                                                                                                     Human; PPP3CB; ss; antisense oligonucleotide; phosphorothioate linkage; 2'-O-methoxyethyl sugar moiety; 5-methylcytosine; autoimmune disorder; Alzheimer's disease; immunosuppressive; nootropic; neuroprotective.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
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                                                                                                                                                                                                                              Human PPP3CB DNA antisense oligonucleotide target region
                                                                                                                                                                                                                                                         06-MAY-2004
                                                                                                                                                                                                                                                                                                              ADJ53600 standard;
                                       31-JUL-2002;
                                                                                                                                              Homo sapiens
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                                                                 31-JUL-2002;
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                                       2002US-00210723
                                                                 2002US-00210723
              PHARM
                                                                                                                                                                                                                                                         (first entry)
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                                                                                                                                                                                                                                                                                                               DNA;
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RESULT 531
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ID ADJ609
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                comprising a 5-methyloytosine. The antisense oligonucleotides are usefu for inhibiting expression of the PPP3CB polypeptide and in preparation a composition for treating autoimmune disorders or Alzheimer's disease. This sequence represents an antisense oligonucleotide target region of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      interleukin; IL-4 receptor; IL-5 receptor; lung disease; airway inflammation; allergy; asthma; impeded respiration; cystic fibrosis; acute respiratory distress syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 4 A; 3 C; 9 G; 4 T; 0 U; 0 Other;
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                                                                                                     Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes (CR1, RANTES, MCP4, useful for prophylaxis or treating respiratory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            05-FEB-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           pulmonary hypertension;
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                                                                                                                                                                                                                          WPI; 2004-203534/19.
                                                                                                                                                                                                                                                                             Nyce JW, Tang L,
Shahabuddin S, Lu
                                                                                                                                                                                                                                                                                                                                                                                                                       29-JUL-2002; 2002US-0399076P
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H, Cong H;
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Claim 2; SEQ ID NO 1809;

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RESULT 532
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    interleukin (IL)-4 receptor, IL-5 receptor or salts of the oligonucleotide and optionally surfactant operatively linked to the
                                                                                                                                                                                                                                                                                                                                                                                          Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes e.g.,
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                                        The present invention relates to an oligonucleotide anti-sense to entitation codon, coding region with 2-10 nucleotides of 5'-end and end of nucleic acid target comprising gene(s) chosen from e.g. interleukin (IL)-4 receptor, IL-5 receptor or salts of the
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Conservative (
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H, Cong H;
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Pred. No.
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                                                                                                                                                           Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory
                                                                                                                                                                                                                                    Nyce JW, Tang
Shahabuddin S,
                                                                                                                                                                                                                                                                                                                                                                                                                              airway inflammation; allergy; asthma; impeded respiration; cystic fibrosis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; oligonucleotide;
                                                                                                                                                   disease e.g.,
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H, Cong H;
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Pred. No.
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The present invention relates to an oligonucleotide anti-sense to e.g., initiation codon, coding region with 2-10 nucleotides of 5'-end and 3'-end of nucleic acid target comprising gene(s) chosen from e.g. interleukin (II) 4 receptor, II-5 receptor or salts of the oligonucleotide and optionally surfactant operatively linked to the oligonucleotide. The method is useful for preventing or treating a respiratory or lung disease, which involves administering to the airways of a subject an effective amount of an inhibitor. The oligonucleotide is useful for production of a medicament for the prevention and/or treatment of a respiratory or lung disease is

Claim 2;

SEQ ID NO 735; 85pp; English.

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RESULT 534
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Matches 20
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interleukin (II)-4 receptor, II-5 receptor or salts of the oligonucleotide and optionally surfactant operatively linked to the oligonucleotide. The method is useful for preventing or treating a respiratory or lung disease, which involves administering to the airways of a subject an effective amount of an inhibitor. The oligonucleotide is useful for production of a medicament for the prevention and/or treatment of a respiratory or lung disease. The respiratory or lung disease is chosen from airway inflammation, allergy(ies), asthma, impeded respiration, cystic fibrosis (CF), chronic obstructive pulmonary diseases (COPD), allergic rhinitis (AR), acute respiratory distress syndrome (ARDS), pulmonary hypertension, lung inflammation, bronchitis, airway obstruction. The present sequence represents an oligonucleotide of the
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Oligonucleotide associated to PDE4C #6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes e.g., CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             06-MAY-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                  end of nucleic acid target comprising gene(s) chosen from e.g. interleukin (IL)-4 receptor, IL-5 receptor or salts of the
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                                                                                                                                                                                                                                                                                                                                                                                                                                   initiation codon,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 2; SEQ ID NO 1796; 85pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            disease e.g., asthma.
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din S, Lu
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                                                                                                                                                                                                                                                                                                                                                                                                                               invention relates to an oligonucleotide anti-sense to codon, coding region with 2-10 nucleotides of 5'-end are
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H, Cong H;
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                                                                                                                                                                                                                                                                                                                                                                                                                                   to e
                                                                                                                                                                                                                                                                                                                                                                                                                                   ω<u>.</u>
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RESULT 535
ADJ60961
ID ADJ609
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SXC
The present invention relates to an oligonucleotide anti-sense to e.g., cc initiation codon, coding region with 2-10 nucleotides of 5'-end and 3'-ecc enterleukin (II)-4 receptor, II-5 receptor or salts of the coligonucleotide and optionally surfactant operatively linked to the coligonucleotide and optionally surfactant operatively linked to the coligonucleotide. The method is useful for preventing or treating a cc respiratory or lung disease, which involves administering to the airways cf a subject an effective amount of an inhibitor. The oligonucleotide is cc useful for production of a medicament for the prevention and/or treatment of a respiratory or lung disease. The respiratory or lung disease is cc chosen from airway inflammation, allergy(ies), asthma, impeded cc respiration, cystic fibrosis (CF), chronic obstructive pulmonary diseases (CODD), allergic rhinitis (AR), acute respiratory distress syndrome cc (ARDS), pulmonary hypertension, lung inflammation, bronchitis, airway construction. The present sequence represents an oligonucleotide of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 interleukin; IL-4 receptor; IL-5 receptor; lung disease;
airway inflammation; allergy; asthma; impeded respiration;
cystic fibrosis; acute respiratory distress syndrome;
pulmonary hypertension; lung inflammation; bronchitis; oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Oligonucleotide associated to PDE4C #27.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20
                                                                                                                                                                                                                                                                                                                                           Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes e.g., CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              06-MAY-2004
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                                                                                                                                                                                                                                                                                                 Claim 2; SEQ ID NO 1817; 85pp; English
                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2004-203534/19
                                                                                                                                                                                                                                                                                                                                                                                                                                              Shahabuddin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          29-JUL-2002; 2002US-0399076P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (EPIG-) EPIGENESIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         643
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                                                                                                                                                                                                                                                                                                                                e.g., asthma.
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din S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CCCAGGCTGGAGTGCAGTGG 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sandrasagra
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Pred. No.
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Query Match

Sequence

20

BP;

4 A;

7 C; 5 G; 4 T; 0 U; 0

Score 20; DB 1; Length 20;

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RESULT 536
ADJ59776
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                δ
                                                                                                                                                    The present invention relates to an oligonucleotide anti-sense to e.g., CC initiation codon, coding region with 2-10 nucleotides of 5'-end and 3'-ec end of nucleic acid target comprising gene(s) chosen from e.g. CC end of nucleotide acid target comprising gene(s) chosen from e.g. CC coligonucleotide and optionally surfactant operatively linked to the coligonucleotide. The method is useful for preventing or treating a CC respiratory or lung disease, which involves administering to the airways co f a subject an effective amount of an inhibitor. The oligonucleotide is CC useful for production of a medicament for the prevention and/or treatment CC aregivatory or lung disease. The respiratory or lung disease is CC chosen from airway inflammation, allergy(ies), asthma, impeded CC respiration, cystic fibrosis (CF), chronic obstructive pulmonary diseases (COPD), allergic rhinitis (AR), acute respiratory distress syndrome CC (ARDS), pulmonary hypertension, lung inflammation, bronchitis, airway CC obstruction. The present sequence represents an oligonucleotide of the
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Matches 20; Conserv
                                                 Matches
                                                              Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  airway inflammation; allergy; asthma; impeded respiration; cystic fibrosis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Oligonucleotide associated to RANTES
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                                                                                                           Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                disease e.g.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2004-203534/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Shahabuddin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           29-JUL-2002; 2002US-0399076P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  interleukin; IL-4 receptor; IL-5 receptor;
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732 AGCTGGGACTACAGGCGCCC
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                                                                Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                  SEQ ID NO 632; 85pp; English.
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in s,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ilarity 100.0%; Pred. No. 1.1e+03; Conservative 0; Mismatches 0;
                                                                                                             BP; 4 A;
                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first
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LuH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                   asthma.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sandrasagra H, Cong H;
                                                              2.0%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 1.1e+03;
                                                                                                             7 C; 7 G; 2 T; 0 U; 0 Other;
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                                                <u>,</u>
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                                                Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  lung disease;
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                                                Indels
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387

CCAAAGTGCTGGGATTACAG

406

0;

Mismatches

Indels

0

Gaps

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                                                                                                                             This sequence represents a primer which was used in the method of the CC invention for diagnosing susceptibility to pereistence or progression of CC cervical intraepithelial neoplasia (CIN) in an individual suffering from CC the disease. The method comprises detecting an allelic deletion in one or CC (progesterone receptor), DLEC1 (deleted in lung and oesophageal cancer 1) CC (progesterone receptor), DLEC1 (deleted in lung and oesophageal cancer 1) CC (progesterone receptor), DLEC1 (deleted in lung and oesophageal cancer 1) CC (progesterone receptor), DLEC1 (deleted in lung and oesophageal cancer 1) CC (progesterone receptor), DLEC1 (deleted in lung and oesophageal cancer 1) CC (progesterone receptor), DLEC1 (deleted in lung and oesophageal cancer 1) CC (progesterone receptor), DLEC1 (deleted in lung and oesophageal cancer 1) CC (progesterone receptor), DLEC1 (deleted from proteins present in the samples CC (derived from non-dyskaryotic and dyskaryotic samples, respectively. The CC (primers, where each pair of primers is suitable for amplifying a CC (primers, where each pair of primers is suitable for amplifying a CC (primers, where each pair of primers is suitable for amplifying a CC (primers) protein selected from FNT3, where each CC (primers) agent is capable of distinguishing between the normal and allelic CC (primers) or DLEC1. The method is useful for diagnosing susceptibility to CC (primers) or protein selected from FNT7, PR, CC (primers) or DLEC1. The method is useful for diagnosing susceptibility to CC (primers) or protein selected from FNT7, PR, CC (primers) or DLEC1. The method is useful for diagnosing susceptibility to CC (primers) or protein selected from FNT7, PR, CC (primers) or DLEC1. The method is useful for diagnosing susceptibility to CC (primers) or DLEC1. The method is useful for diagnosing susceptibility to CC (primers) or DLEC1. The method is useful for diagnosing susceptibility to CC (primers) or DLEC1.
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Query Match
Best Local S
Matches 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ss; primer; diagnosis; cervical intraepithelial neoplasia; CIN; allelic deletion; FHIT; fragile histidine triad gene; PR; progesterone receptor; DLEC1; deleted in lung and oesophageal cancer 1; TRIM29; tripartite motif-containing 29; microsatellite; D3S1300; D3S1260; D11S35; D11S528.
                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; SEQ ID NO 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      present in the samples.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Diagnosing cervical intraepithelial neoplasia comprising detecting an allelic deletion in genes selected from FHIT, PR, DLEC1- or TRIM 29 by comparing the FHIT, PR, DLEC1 and/or TRIM 29 polynucleotides or proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  24-AUG-2002;
26-AUG-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   04-MAR-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Primer #1 for amplification of D6S105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADL23339;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20-AUG-2003; 2003WO-GB003637.
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                             Similarity
                                                                                              20 BP; 6
  2.0%;
llarity 100.0%;
Conservative
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2002US-0405717P
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                                                                                              A; 4 C; 6 G; 4 T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        56pp;
                             Score 20;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        English.
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                                                  DB 1;
                             1.1e+03;
                                                                                                0 Other;
                                                Length 20;
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by
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RESULT 539
ADL61592/c
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ADL32388/c
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 Query Match
Best Local
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11-JAN-2000;
02-MAY-2000;
07-JUL-2000;
                                                                                                                                                                                                                                                                      This invention relates to a novel primers useful for synthesising full length cDNA molecules that encode human proteins. Specifically, it refers to secretory or membrane proteins that are potential therapeutic agents/ target molecules in the field of medicine, and in particular genes encoding proteins that are associated with signal transduction, allycoproteins and transcription. The present invention describes a method for efficiently cloning a full length human cDNA from both the 5' and 3' ends using the oligo-capping method. This oligonucleotide sequence is a human clone specific PCR primer used in an exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADL32388 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 human; medicine; signal transduction; glycoprotein; transcription; oligo-capping method; ss; PCR; primer.
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                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                     New oligonucleotide primers (830 cDNAs) useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Wakamatsu
                                                    03-JUN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (REAS-)
predictor set; protein tyrosine kinase
antiangiogenic; vasotropic; vulnerary;
                               Human
                                                                                                ADL61592
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                                                                                                                                                                          388
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                               protein
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                                                                                                                                                                                                                                                                                                                                                                                                            human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nishikawa T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RES ASSOC BIOTECHNOLOGY
                                                                                                                                                                                                          Similarity
                                                                                                                                                                          CAAAGTGCTGGGATTACAGG 407
                                                                                                                                                                                                                                          20
                                                                                               standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; 99JP-00194486.
; 2000JP-00118774.
; 2000JP-00183865.
; 2000EP-00114089.
                                                                                                                                                                                                                                           BP; 4 A;
                                                                                                                                                                                                                                                                                                                                                                                      SEQ ID
                                                                                                                                                                                               2.0%;
llarity 100.0%;
Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                            CDNAs.
                                                    (first entry)
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                                tyrosine kinase biomarker-related RT-PCR primer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   primer to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA;
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                                                                                                DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        entry)
                                                                                                                                                                                                                                          7 C; 3 G;
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na T, Nagai
                                                                                                                                                                                                                                                                                                                                                                                      4421; 1340pp;
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                                                                                                                                                                                                          Score 20;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   amplify human full length cDNA SeqID 4421.
                                                                                                                                                                                                                                           6 T;
                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hayashi K,
K, Kojima
                                                                                                                                                                                                                                           0 U;
                                                                                                                                                                                                          No. 1.1e+03;
                                                                                                                                                                                                                                                                                                                                                                                      English.
biomarker; cytostatic;
pharmacogenomic; drug
                                                                                                                                                                                                                      DB 1;
                                                                                                                                                                                                                                           0 Other;
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S, Otsuki
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                                                                                                                                                                                                                    Length 20;
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T, Ko
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RESULT 540
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Best Local S
Matches 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      polynucleotides and/or polypeptides whose expression pattern is predictive of the response of cells to treatment with a compound that modulates protein tyrosine kinase activity or members of the protein tyrosine kinase pathway. The molecules of the invention demonstrate cyrosine kinase pathway. The molecules of the invention demonstrate cyrostatic, antiangiogenic, vasotropic and vulnerary activities and may be useful in the field of pharmacogenomics, in particular for determining drug sensitivity and in treating breast cancer, hypervascular diseases, anglogenesis and scars in wound healing. The current sequence is that of a human protein tyrosine kinase biomarker-related RT-PCR primer of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New predictor sets with a plurality of polynucleotides and/or polypeptides whose expression pattern predicts cell response to compound that modulates protein tyrosine kinase activity, useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        breast cancer; hypervascular disease; angiogenesis; wound healing human; ss; RT-PCR; PCR; primer.
                                                                                                chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1; motion cardiant; neuroprotective; antiinflammatory; neuroprotective; antiinflammatory; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; SEQ ID NO 516; 649pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    treating breast cancer.
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immunomodulatory; cardiovascular; gene therapy; inflammation Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorcardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a novel predictor set comprising a plurality
                                                                                                                                                                                                                                                        Human mPGES-1 chimeric antisense oligonucleotide SEQ
                                                                                                                                                                                                                                                                                                             01-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                ADM14394;
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Pred. No.
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                                                                             gene therapy; inflammation;
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RESULT 541
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Best Local S
Matches 20
                                                                                                                                                                                                                                                                                      mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, antiarchritic, vasotropic, antihiflammatory, neuroprotective, nootropic, antiarchritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
01-JUL-2004
                              ADM14746
                                                            ADM14746 standard; DNA; 20
                                                                                                                                                                                                                                                              Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9034.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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                                                                                                                                                                                                                 Similarity
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                                                                                                                                                                    GGCCTCCCAAAGTGCTGGGA 870
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(first entry)
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antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, antiinflammatory, neuroprotective, nootropic, antianthritic, vasotropic, antiinflammatory, neuroprotective, nootropic, antianthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or
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                                                                                                                                                                                               The present sequence represents a chimeric antisense oligonucleotide targeted to human microscomal prostaglandin E2 synthase (mpGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9q34.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric
                                                                                                                                                                                                                                                                                                                                                                                                                                             New antisense compound, having a sequence targeted to a nucleic acencoding mPGES-1, useful for preparing a composition for treating inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gierse JK
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                                    ophthalmic,
                                                                                                                                                                                                                                                                                                                                                                                         Claim 4; SEQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                 ischemia.
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                                  immunorogicai,
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/note= "2'.-O-methocyethyls"
16. .20
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/mod_base=
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'note= "2'-O-methoxyethyls"
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                                    cardiovascular
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                                  or neurological
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     a nucleic acid
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                                      disorder
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Sequence

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ADM14277/c
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cardiovascular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:464
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9q34.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding
                                                                                                                New antisense compound, having a sequence targeted to a nucleic acencoding mPGES-1, useful for preparing a composition for treating inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                               modified_base
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                                                                           SEQ ID NO 464; 132pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         antisense oligonucleotide; phosphorothicate;
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ar disorder;
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J. 1.1e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                   chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mpGES-1 imPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; notropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric
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                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:669.
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16. .20
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== "2'-O-methoxyethyls"
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_(PHAA) PHARMACIA CORP

25-SEP-2002; 2002US-0413549P 25-SEP-2003; 2003WO-US030374

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
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/*tag= a
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residues are 5-methylcytidines"
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               chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; mPGES-1; mCGES-1; motidiabetic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; antiinflammatory; neuroprotective; antiarthritic; vasotropic; ophthalmological;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to 9q34.3. The present invention also describes: (1) antisense compounds,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New antisense compound, having a sequence targeted to encoding mPGES-1, useful for preparing a composition f
                                                                                                                                                     01-JUL-2004
                                                                                                                                                                                                                     ADM15160
                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 4; SEQ ID NO 1496; 132pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2004-305094/28
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                                                                                                                    Human
                                                                                                                                                                                    ADM15160;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         encoding mPGES-1, useful for preparing a composition for treating inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (PHAA ) PHARMACIA CORP
 mmunomodulatory; cardiovascular;
                                                                                                                                                                                                                                                                                                                                     769
                                                                                                                  mPGES-1
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                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                     standard; DNA;
                                                                                                                                                                                                                                                                                                                      TTTTGTATTTTTAGTAGAG 788
                                                                                                                                                                                                                                                                                                     TTTTTGTATTTTAGTAGAG
                                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 12
                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                   (first entry)
                                                                                                                chimeric antisense oligonucleotide SEQ
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/note= "2'-O-methoxyethyls"
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                                                                                                                                                                                                                                                                                                                                                                                                                                       A; 4 C; 0 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                      100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                       2.0%;
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                                                                                                                                                                                                                                                                                                                                                                                      Score 20;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                      Mismatches
gene
therapy;
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                                                                                                                                                                                                                                                                                                                                                                                                      DB 1;
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                                                                                                                    NO:1347
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                                                                          Query Match
Best Local
                                                              Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 modified_base
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Synthetic.
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                                                                                                                                                               antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.ginflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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                                                                                                                       Sequence
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 20
                                                                          Similarity
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                   CCTCGGCCTCCCAAAGTGCT 866
 CCTCGGCCTCCCAAAGTGCT
                                                                                                                       BP; 4 A;
                                                                                                                                                    immunological, cardiovascular or neurological
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16. .20
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/mod_ba
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/mod_base= OTHER
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                                                                          2.0%;
                                                                                                                       4 C; 9 G; 3 T; 0 U; 0 Other;
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                                                                          Score 20;
Pred. No.
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                                                              Mismatches
                                                                                           DB 1;
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                                                                           .1e+03;
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RESULT 546
ADM14957/c
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                                                                                                                           The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPCES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9334.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cycostatic, antisense oligonucleotides and antisense compounds have cycostatic.
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Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.ginflammation, Alzheimer's disease, arthritis, diabetes, cancer or
antidiabetic, immunomodulator, cardiant, neuroprotective, antiarthritic, vasotropic, antiarthritic, vasotropic, antiarthritic, immunomodulatory and cardiovascular activities, and can ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mpgBS-1 inhibitors and in gene therapy. The antisense compound
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/mod_base= OTHER
/mote= "2'-O-methoxyethyls"
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/mod_base= OTHER
/note= "2'-O-methocyethyls"
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ADM15553/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; antiarthritic; vasotropic; ophthalmological; immunomodulator; nootropic; antiarthritic; vasotropic; ophthalmological;
                   New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating entirely inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                  Gierse
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human mPGES-1
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           ischemia
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                                                                                                                                                                            25-SEP-2003; 2003WO-US030374.
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                                                                                                                          (PHAA ) PHARMACIA CORP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   chimeric antisense oligonucleotide SEQ ID NO:1740.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     disorder; neurological disorder; ss.
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                                                                                                                                                                                                                                                                                                                                                         /note= "phosphorothioate linkages and
residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                               *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA;
                                                                                                                                                                                                                                                                                                                                                                                  _base= OTHER
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                                                                                                                                                                                                                                                                                                       "2'-0-methocyethyls"
                                                                                                                                                                                                                                                       "2'-O-methoxyethyls"
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; Pred. No.
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hes 0;
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RESULT 548
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    934.3. The present invention also describes: (1) antisense compounds, 934.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mpGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mpGES-1. MpGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antisense oligonucleotides and antisense compounds have cytostatic, antisense oligonucleotides, cardiant, neuroprotective, cardiant, neuroprotective, on antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, isohaemia or reperfusion injury, or
                                                                                                                                                                                                                                                                                                                                                                                                  immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological;
                                                                                                                                                                                                                                                                                                                                             Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 4; SEQ
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                                                                                                                       modified_base
                                                                                                                                                                                             modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              chimeric; antisense oligonucleotide; phosphorothioate; human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:1268
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sequence represents a chimeric antisense oligonucleotide human microsomal prostaglandin E2 synthase (mPGES-1). The
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/mod_base=
                                                                                                                                                                                                                  residues
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                                                                      'mod_base= OTHER
'note= "2'-O-methoxyethyls"
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Pred. No.
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08-APR-2004.

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ARBSULT 549
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Best Local 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9934.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           antidiabetic, immunomodulator, cardiant, neuroprotective, vasotropic, antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis diabetes, cancer, ischaemia or reperfusion injury, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2004-305094/28
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    25-SEP-2002; 2002US-0413549P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          antisense oligonucleotides and antisense compounds have cytostatic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 4; SEQ ID NO 1268; 132pp; English.
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                                                                                                Homo sapiens.
Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:1455
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modified_base
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                                                                                                                                                                                                      disorder;
   Location/Qualifiers
1. .20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA;
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Pred. No.
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BXXXXXX

ADM14958

standard;

DNA;

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01-JUL-2004

(first entry)

Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:1145.

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RESULT 550
ADM14958/c
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                                                                                                                                                                                                               Query Match
Best Local S
Matches 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         modified_base
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                                                                                                                                                                                                                                                                                                                               BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   human microsomal prostaglandin E2 synthase (mPGES-1).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sequence represents a chimeric antisense oligonucleotide
                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                               3 A;
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== "2'-O-methoxyethyls"
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                                                                                                                                                                                                                     0;
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Pred. No.
                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                    1.1e+03;
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chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mpEES-1; mpEES-1 inhibitor; microsomal prostaglandin E2 synthase; mpEES-1; mpEES-1 inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                   The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9934.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antisense oligonucleotides and antisense compounds have cytostatic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g. inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              25-SEP-2003; 2003WO-US030374.
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                                                                   antidiabetic, immunomodulator, cardiant, neuroprotective, association, neuroprotective, neuroprotective, nootropic, antiarrhritic, vasotropic, antiarrhritic, vasotropic, antiarrhritic, vasotropic, antiarrhritic, vasotropic, antiarrhritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's
                                                                                                                                                                                                                                                                                                                                                                                                              Claim 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25-SEP-2002; 2002US-0413549P
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                                                                                                                                                                                                                                                                                                                                                                                                              SEQ ID NO 1145; 132pp; English
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Query Match Best Local Similarity

100.0%; 2.0%;

Score 20; Pred. No.

1.1e+03; DB 1; 0 Other;

Length

20

Sequence 20

BP; 4

A; 5 C; 8 G; 3 T; 0 U;

arthritis,

diabetes,

immunological,

cardiovascular

cancer,

ischaemia or reperfusion injury,

õ

neurological

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The invention relates to oligonucleotides anti-sense to an initiation codon, coding region, 5' or 3' intron-exon junction, intron or region with 2-10 nucleotides of the 5'-end or 3'-end of a nucleic acid target chosen from a gene encoding interleukin (II) -4 receptor, interleukin (II) compared to the from a gene encoding interleukin (II) -4 receptor, interleukin (II) compared to the first place by PDE4 A, PDE4 B, PDE4 C or PDE4 D. The invention compared to a method of screening a candidate compound that binds to one or more nucleic acid target(s) or expressed product(s), for the compound the first place and the first place are useful for reducing or inhibiting expression of a conjunctation and/or treatment of a respiratory or lung disease. The compound the first place are considered and compound the first place are considered and compound the first place are considered as a compound the first place and considered and compound the first place are considered as a compound the first place are considered as a compound the first place are considered as a compound the first place and compound the first place are considered as a compound the first place and compo
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                                                                                                                                                                                                                                                                                                                                                                                                           Novel single or multiple initiation codon, intron RANTES, MCP4, useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nусе JW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (NYCE/)
(SAND/)
(TANG/)
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                                                                                                                                                                                                                                                                                                                                              Claim 2;
                                                                                                                                                                                                                                                                                                                                                                                             asthma.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 inflammation; bronchitis; airway obstruction;
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CCR3; Eotaxin-1; RANTES; MCP4; CD23; ICAM; VCAM; tryptase a;
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                                                                                                                                                                                                                                                                                                                                                SEQ ID
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tang L,
ong H;
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Sequence 20

BP; 3 A;

4 C; 10

<u>ن</u>

3 T; 0 U; 0 Other;

DB 1;

Length 20

respiratory or lung disease is associated with hyper-responsiveness to and/or increased levels of, adenosine and/or levels of adenosine A receptor(s), and/or asthma and/or lung allergies associated with inflammation or an inflammatory disease. The respiratory or lung disease is chosen from airway inflammation, allergy, asthma, impeded respiration, cystic fibrosis (CP), chronic obstructive pulmonary disease (COPD), allergic rhinitis, acute respiratory distress syndrome, pulmonary hypertension, lung inflammation, bronchitis, airway obstruction or bronchoconstriction. This sequence represents an oligonucleotide of the

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RESULT 552
ADO46429
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Best Local S
Matches 20
                                                                                                                                                                                                                                                                                                                                                                                          Human; ss; interleukin-4 receptor; IL-4; interleukin-5 receptor; IL-5; CCR1; CCR3; Botaxin-1; RANTES; MCP4; CD23; ICAM; VCAM; tryptase a; tryptase b; PDE4 A; PDE4 B; PDE4 C; PDE4 D; respiratory disease; lung disease; hyper-responsiveness; adenosine; adenosine A receptor; asthma; lung allergy; inflammation; inflammatory disease; airway inflammation; allergy; impeded respiration; cystic fibrosis; CF; chronic obstructive pulmonary disease; COPD; allergic rhinitis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; airway obstruction; bronchoconstriction.
                                                                                                                                                                                                      (NYCE/)
(SAND/)
(TANG/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human oligonucleotide #1795
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23-APR-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADO46429 standard;
                                 Novel single or multiple target oligonucleotide anti-sense to e.g. Coinitiation codon, intron of respiratory disease-relevant gene e.g. Corantes, MCP4, useful for prophylaxis or treating respiratory disease
                                                                                                         Nyce JW, Sandrasagra A, Ta
Shahabuddin S, Lu H, Cong
                                                                                                                                                                                                                                                                                            25-JUL-2003; 2003US-00627930.
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Claim 2;
                                                                                   WPI; 2004-293804/27
                                                                                                                                                                     (SHAH/)
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                                                                                                                                                                  MILLER S.
SHAHABUDDIN S.
                                                                                                                                                                                                        SANDRASAGRA
TANG L.
                                                                                                                                                                                            AGUILAR D.
SEQ
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llarity 100.0%;
Conservative
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2002WO-US013143.
ID NO 1796; 174pp; English
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                                                                                                          Tang L, ong H;
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                                                                                                                                                                                                                                          The invention relates to oligonucleotides anti-sense to an initiation codon, coding region, 5' or 3' intron-exon junction, intron or region codon, coding region, 5' or 3' intron-exon junction, intron or region codon, coding region, 5' end or 3'-end of a nucleic acid target chosen from a gene encoding interleukin (II)-4 receptor, interleukin (IL)-5 receptor, CCR1, CCR3, Ectaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, CC tryptase a, tryptase b, PDE4 A, PDE4 B, PDE4 C PDE4 D. The invention CC also relates to a method of screening a candidate compound that binds to come or more nucleic acid target(s) or expressed product(s), for the come or more nucleic acid target(s) or expressed product(s), for the come or more nucleic acid target(s) or expressed product(s), for the colligonucleotides are useful for reducing or inhibiting expression of a CC gene or mRNA encoding interleukin-4 receptor, interleukin-5 receptor, CC CCR1, CCR3, Ectaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, CC tryptase b, PDE4 A, PDE4 B, PDE4 C, or PDE4 D. The oligonucleotides are CC tryptase b, PDE4 A, PDE4 B, PDE4 C, or PDE4 D. The oligonucleotides are CC tryptase b, PDE4 A, PDE4 B, PDE4 C, or PDE4 D. The oligonucleotides are CC tryptase b, and/or sthma and/or lovels of adenosine A CC and/or increased levels of, adenosine and/or levels of adenosine A CC and/or increased levels of, adenosine and/or levels of adenosine A CC and/or increased levels of, adenosine and/or levels of adenosine A CC and/or increased levels of, adenosine and/or levels of adenosine A CC and/or increased levels of, adenosine and/or levels of adenosine A CC and/or increased levels of, adenosine and/or levels of adenosine A CC and/or increased levels of, adenosine and/or levels of adenosine A CC and/or increased levels of, adenosine and/or levels of adenosine (COPD), consite of the constructive pulmonary disease (COPD), consite of the constructive pulmonary disease (COPD)
Sequence 20 BP; 3 A; 5 C; 9 G; 3 T; 0 U; 0 Other;
                                                                                                                                  allergic rhinitis, acute respiratory distress syndrome, pulmonary hypertension, lung inflammation, bronchitis, airway obstruction or bronchoconstriction. This sequence represents an oligonucleotide of the
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Query Match Best Local Matches 643 CCCAGGCTGGAGTGCAGTGG 662 20; Similarity Conservative 2.0%; 0, Score 20; Pred. No. 20 Mismatches DB 1; Length 20; 1.1e+03; Indels 0 Gaps

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RESULT 553 ADO46442 ID ADO464 XX lung disease; hyper-responsiveness; adenosine; adenosine A receptor; asthma; lung allergy; inflammation; inflammatory disease; airway inflammation; allergy; impeded respiration; cystic fibrosis; chronic obstructive pulmonary disease; COPD; allergic rhinitis; acute respiratory distress syndrome; pulmonary hypertension; Human; ss; interleukin-4 receptor; IL-4; interleukin-5 receptor; IL-5; CCR1; CCR3; Eotaxin-1; RANTES; MCP4; CD23; ICAM; VCAM; tryptase a; tryptase b; PDE4 A; PDE4 B; PDE4 C; PDE4 D; respiratory disease; ADO46442 standard; DNA; 20 BP Human oligonucleotide #1808 15-JUL-2004 (first entry)

(SAND/)
(TANG/) 23-APR-2002; 23-APR-2002; 25-JUL-2003; 11-MAR-2004. US2004049022-A1 (NYCE/) TANG SANDRASAGRA NYCE J W 2002WO-US013135 2002WO-US013143 2003US-00627930

Homo sapiens

inflammation; bronchitis;

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Best Local Similarity
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                                                                              Human, 88; interleukin-4 receptor; II-4; interleukin-5 receptor; II-5; CCR1; CCR3; Eotaxin-1; RANTES; MCP4; CD23; ICAM; VCAM; tryptase a; tryptase b; PDE4 A; PDE4 B; PDE4 C; PDE4 D; respiratory disease; lung disease; hyper-responsiveness; adenosine; adenosine A receptor; asthma; lung allergy; inflammation; inflammatory disease; airway inflammation; allergy; impeded respiration; cystic fibrosis; cfronic obstructive pulmonary disease; COPD; allergic rhinitis;
                                                                                                                                                                                                                                                                        Human oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                          ADO45266 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20
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                                                             acute respiratory distress syndrome; pulmonary hypertension
                                                                                                                                                                                                                                                                                                                     15-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             bronchoconstriction. This sequence represents an oligonucleotide of the
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                                        bronchitis; airway obstruction; bronchoconstriction
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RESULT 555 ADO46449 ID ADO464

ADO46449 standard; DNA; 20

ВP

ADO46449

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732 1

AGCTGGGACTACAGGCGCCC 751

20

Query Match Best Local S Matches 20

Similarity

2.0%; Score 20; DB 100.0%; Pred. No. 1. Live 0; Mismatches

DB 1; L 1.1e+03;

Length 20;

0;

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Gaps

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Conservative

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The invention relates to oligonucleotides anti-sense to an initiation codon, coding region, 5' or 3' intron-exon junction, intron or region with 2-10 nucleotides of the 5'-end or 3'-end of a nucleic acid target chosen from a gene encoding interleukin (II)-4 receptor, interleukin (II) -5 receptor, CCR1, CCR3, Botaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, CC tryptase a, tryptase b, PDE4 A, PDE4 B, PDE4 C or PDE4 D. The invention CC also relates to a method of screening a candidate compound that binds to one or more nucleic acid target(s) or expressed product(s), for the CC one or more nucleic acid target(s) or expressed product(s), for the CC one or more nucleic acid target(s) or expressed product(s), for the CC oligonucleotides are useful for reducing or inhibiting expression of a gene or mRNA encoding interleukin-4 receptor, interleukin-5 receptor, CC CCR1, CCR3, Botaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, CC tryptase b, PDE4 A, PDE4 B, PDE4 C, or PDE4 D. The oligonucleotides are cuseful for preventing or treating a respiratory or lung disease. The CC useful for preventing or treating a respiratory or lung disease. The CC useful for preventing or treating a respiratory or lung disease. The CC cand/or increased levels of, adenosine and/or levels of adenosine A conceptor(s), and/or asthma and/or lung allergies associated with CC inflammation or an inflammatory disease. The respiratory or lung disease is chosen from airway inflammation, allergy, asthma, impeded respiration, CC allergic rhinitis, acute respiratory distress syndrome, pulmonary or increased (CPP), chronic obstructive pulmonary disease (COPD), allergic rhinitis, acute respiratory distress syndrome, pulmonary or lung disease.
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Sequence
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23-APR-2002;
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15-JUL-2004

(first

entry)

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chosen from a gene encoding interleukin (IL)-4 receptor, interleukin (IL)

CC chosen from a gene encoding interleukin (IL)-4 receptor, interleukin (IL)

CC receptor, CCR1, CCR3, Ectaxin-1, RANTES, MCP4, CD23, ICAM, VCAM,

CC tryptase a, tryptase b, PDE4 A, PDE4 B, PDE4 C or PDE4 D. The invention

CC also relates to a method of screening a candidate compound that binds to

CC one or more nucleic acid target(s) or expressed product(s), for the

CC one or more nucleic acid target(s) or expressed product(s), for the

CC oligonucleotides are useful for reducing or inhibiting expression of a

CC oligonucleotides are useful for reducing or inhibiting expression of a

CC oligonucleotides are useful for reducing or inhibiting expression of a

CC gene or mRNA encoding interleukin-4 receptor, interleukin-5 receptor,

CC CCR1, CCR3, Ectaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a

CC tryptase b, PDE4 B, PDE4 C, or PDE4 D. The oligonucleotides are

CC useful for preventing or treating a respiratory or lung disease. The

CC useful for preventing or treating a respiratory or lung disease. The

CC and/or increased levels of, adenosine and/or levels of adenosine A

CC receptor(s), and/or asthma and/or lung allergies associated with

CC inflammation or an inflammatory disease. The respiratory or lung disease

CC office therefore a company inflammation, allergy, asthma, impeded respiration,

CC crystic fibrosis (CF), chronic obstructive pulmonary disease (COPD),
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codon, intron of respiratory disease-relevant gene e.g. CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g.
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23-APR-2002; 2002WO-US013143.
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allergic rhinitis, acute respiratory distress syndrome, pulmonary hypertension, lung inflammation, bronchitis, airway obstruction or bronchoconstriction. This sequence represents an oligonucleotide of the

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RESULT 556
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Best Local Sim
Matches 20;
The invention relates to oligonucleotides anti-sense to an initiation codon, coding region, 5' or 3' intron-exon junction, intron or region with 2-10 nucleotides of the 5'-end or 3'-end of a nucleic acid target chosen from a gene encoding interleukin (IL)-4 receptor, interleukin (II -5 receptor, CCR1, CCR3, Eotaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, tryptase b, PD24 A, PD24 B, PD24 C or PD24 D. The invention also relates to a method of screening a candidate compound that binds to one or more nucleic acid target(s) or expressed product(s), for the prevention and/or treatment of a respiratory or lung disease. The oligonucleotides are useful for reducing or inhibiting expression of a gene or mRNA encoding interleukin-4 receptor, interleukin-5 receptor, CCR1, CCR3, Eotaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     asthma; lung allergy; inflammation; inflammatory disease; airway inflammation; allergy; impeded respiration; cystic fibrosis; CF; chronic obstructive pulmonary disease; COPD; allergic rhinitis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; airway obstruction; bronchoconstriction.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; ss; interleukin-4 receptor; IL-4; interleukin-5 receptor; IL-5; CCR1; CCR3; Ectaxin-1; RANTES; MCP4; CD23; ICAM; VCAM; tryptase a; tryptase b; PDE4 A; PDE4 B; PDE4 C; PDE4 D; respiratory disease; lung disease; hyper-responsiveness; adenosine; adenosine A receptor; asthma; lung allergy; inflammation; inflammatory disease;
                                                                                                                                                                                                                                            Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codon, intron of respiratory disease-relevant gene e.g. CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g.
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CONG H.
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in S, Lu H, Cong
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                      The invention describes a method of typing (M1) a gene (I) that has one or more polymorphic microsatellite loci (PML). The method comprises: PCR amplification of at least one DNA region of (I) that includes PML, using as template a DNA sample containing at least one segment of (I); and determining the length of the resulting amplicon(s). Also described are: a method of determining (M2) microsatellite markers (MM) for predisposition to a disease, associated with a gene that includes one or more PML; and prediagnosis (M3) of diseases associated with gene that include PML. The method is used to identify microsatellite markers, in a disease-related gene, that are associated with a predisposition to
                                                                                                                                                                                                                                                                                                                                                                                                                                             Typing genes that contain polymorphic microsatellite loci, useful for identifying predisposition to disease, by amplification and determining length of amplicons.
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      prediagnosis of such diseases,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      cytostatic; gene therapy; inhibitors of apoptosis-like; IAP-like; IAP-like modulator; IAP-like associated disorder; hyperproliferative disorder; human; antisense oligonucleotide; antisense technology; ss.
                                                  New compound targeted to a nucleic acid molecule encoding apoptosis (IAP)-like and inhibits expression of IAP-like, modulating the expression of IAP-like or for treating, e.g hyperproliferative disorder.
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/note= "OTHER= 2'-O-Methoxyethyl
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- "OTHER= 2'-O-Methoxyethyl
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 1;
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                                                                                                                                                                                                                                                                                                                                                                    backbone. All cytidines
                                                                                                                                                                                                                                                                    (2'-MOE)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                               -MOE)
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                                                                           j inhibitors
useful for
                                                                                                                                                                                                                                                                    nucleotides"
                                                                                                                                                                                                                                                                                                               nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         quicker and
equencing. This
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             human prion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                       of.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0,
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The invention

describes

a compound 8-80 nucleobases in length targeted encoding inhibitors of apoptosis (IAP)-like,

ç

nucleic acid molecule

Example 14; SEQ ID

NO 83;

58pp; English

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RESULT 559
AD052273/c
ID AD0522
XX AD0522
XX AD0522
XX AD0522
XX IAP-li
KW IAP-li
KW IAP-li
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KW hypert
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Best Local S
Matches 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cytostatic; gene therapy; inhibitors of apoptosis-like; IAP-like; IAP-like modulator; IAP-like associated disorder; hyperproliferative disorder; human; antisense oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human inhibitor of apoptosis-like antisense oligonucleotide seqid 149.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADO52273 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 oligonucleotide.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  antisense technology; ss.
                                                                                                                                                                                                                                                 22-NOV-2002;
                                                                                                                                                                                                                                                                                                           22-NOV-2002; 2002US-00303325
                                                                                                                          Bennett
                                                                                                                                                                                       (ISIS-) ISIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  969 CTCGGCTCACTGCAACCTCT 988
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                                                                                                                          CF,
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nilarity 100.0%;
Conservative 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                 2002US-00303325
                                                                                                                          Dobie
                                                                                                                                                                                          PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note= "
15. .20
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/note= "OTHER= Phosphorothioate backbone.
are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /mod_base= OTHER
/note= "OTHER= 2'-O-Methoxyethyl (2'-MOE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               _base= OTHER
e= "OTHER= 2'-O-Methoxyethyl (2'-MOE)
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Pred. No.
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1.1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      nucleotides'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       All cytidines
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New compound

targeted to

a nucleic acid molecule encoding inhibitors

e E

The primers AAV27964-V28086 are used to amplify ataxia telangiectasia (ATM) exons and their adjacent splice junction sites. These can be use as a method of detecting a mutation in the ATM gene by comparing the

PCR

Claim 6; Page 6; 47pp; English.

WPI; 2004-399725/37.

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RESULT 560
AAV27991/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention describes a compound 8-80 nucleobases in length targeted to compound encoding inhibitors of apoptosis (IAP)-like, compound specifically hybridises with the nucleic acid molecule encoding inhibitors of apoptosis (IAP)-like.

CC where the compound specifically hybridises with the nucleic acid molecule encoding IAP-like comprising 16000 bp (SEQ ID NO. 4) and inhibits the compound in the sequence comprision of the presence of IAP-like. Also described are: inhibiting the expression of EAP-like in a sample using state comprising identifying the presence of IAP-like in a sample using at least one of the primers comprising SEQ ID NO. 5 or 6, or the probe comprising SEQ ID NO. 7; a kit or assay device comprising the compound; and treating an animal having a disease or condition associated with IAP-CC and treating an animal having a disease or condition associated with IAP-CC it is also useful for diagnosing or treating diseases associated with CC expression of IAP-like, e.g. a hyperproliferative disorder. This sequence comprisors human inhibitor of apoptosis (IAP)-like antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local S
Matches 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            apoptosis (IAP)-like and inhibits expression of IAP-like, useful modulating the expression of IAP-like or for treating, e.g. hyperproliferative disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 5 A; 3 C; 9 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 14; SEQ ID NO 147; 58pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                             ss; PCR; primer; amplification; ataxia telangiectasia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ataxia telangiectasia exon 17 primer 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25-SEP-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAV27991 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    oligonucleotide.
                                                                                                       Method of detecting ataxia telangiectasia - based on intron-exon boundaries, useful for
                                                                                                                                                                                                                                                                          17-NOV-1997;
                                                                                                                                                                                                                                                                                                                                    WO9822621-A1
                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAV27991;
                                                                                                                                                  WPI; 1998-312503/27
                                                                                                                                                                                 Concannon
                                                                                                                                                                                                              (VIRG-) VIRGINIA MASON RES CENT
                                                                                                                                                                                                                                           20-NOV-1996;
                                                                                                                                                                                                                                                                                                       28-MAY-1998
                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                radiation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           969 CTCGGCTCACTGCAACCTCT 988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20
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                                                                                                                                                                                                                                                                                                                                                                                                                  breast
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2.0%; Sullarity 100.0%; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                            96US-00753147.
                                                                                                                                                                                                                                                                          97WO-US020953
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1.1e+03;
                                                                                                       comprises use of diagnosing diseas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 20;
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                                                                                                          use of primers
disease in
                                                                                                                                                                                                                                                                                                                                                                                                                                 diagnosis; human;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 561
                            The present invention describes a restriction primer for eukaryotic short interspersed repeated sequences (SINE), which has one or more additional bases that are a mismatch to, or are unrelated to, the 3'-terminal end of the SINE. The annealing temperature of the primer to the NNA sequence is kept higher than the fusion temperature of the primer during polymerase chain reaction (PCR). The PCR fragments obtained are subjected to electrophoresis to obtain a fingerprint. By comparing the polymorphs from the electrophoresis band pattern, eukaryotic individuals are distinguished. The primer is used for amplifying a eukaryotic distinguished. The primer is used for amplifying a eukaryotic decoxyribonucleic acid (DNA) sequence, pinched between SINE sequences by polymerase chain reaction (PCR) fingerprinting. In particular it may be used individual identification of humans for medical and legal applications and ecological studies. NNA specimens in traces (approximately 10 ng in mass) can be used for individual discrimination of eukaryotes using the primer in a polymerase chain reaction (PCR).

ANZESI43 to ANZESI91 represent specifically claimed examples of primers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   products of amplification from a sample from a patient suspected of having an ATM mutation with a sample from a non-mutated ATM patient This method is especially useful for diagnosing ataxia telangiectasia in heterozygotes and can be used to locate the positions of the mutation. The diagnosis of ataxia telangiectasia in patients needing therapeutic radiation will prevent fatal radiation burns and the development of
                                                                                                                                                                                                                                                                                                                                                                                                      Restriction primer for distinguishing individuals with short interspersed repeated sequence of eukaryotes by restricted polymerase chain reaction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 JP2913035-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Oncorhynchus; restriction primer; short interspersed repeated sequence; eukaryote; restricted polymerase chain reaction fingerprinting; identification; DNA specimen; discrimination; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 21
                                                                                                                                                                                                                                                                                                                                                   Claim 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1999-583348/50.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        10-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human short interspersed repetitive element
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      13-DEC-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAZ25145
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; short interspersed repetitive element; SINE; PCR; primer;
                                                                                                                                                                                                                                                                                                                                                                                       fingerprinting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NORINSUISANSHO SUISANCHO YOSHOKU KENKYUSHOCHO
                                                                                                                                                                                                                                                                                                                                                   Page 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CCCAGGCTGGAGTGCAGTGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ilarity 100.0%;
Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP; 3 A; 10 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    98JP-00195692
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          98JP-00195692
                                                                                                                                                                                                                                                                                                                                                 17pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2.0%;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      N
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length
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868

GGATTACAGGCGTGAGCCAC 887

Matches Query Match Best Local :

20;

Conservative

0

Mismatches

Similarity

2.0%; 5 C; 7

Score 20; Pred. No. 3 T;

1.2e+03;

Indels

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Gaps

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Length 21;

Sequence

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U; 0 Other; DB 1;

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ARESULT 562
ARZZ513
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                                                                                The present invention describes a restriction primer for eukaryotic short CC interspersed repeated sequences (SINE), which has one or more additional CC bases that are a mismatch to, or are unrelated to, the 3'-terminal end of CC the SINE. The annealing temperature of the primer to the DNA sequence is kept higher than the fusion temperature of the primer during polymerase CC chain reaction (PCR). The PCR fragments obtained are subjected to electrophoresis to obtain a fingerprint. By comparing the polymorphs from CC the electrophoresis band pattern, eukaryotic individuals are distinguished. The primer is used for amplifying a eukaryotic deoxyribonucleic acid (DNA) sequence, pinched between SINE sequences by CC oblymerase chain reaction (PCR) fingerprinting. In particular it may be used individual identification of humans for medical and legal capplications and ecological studies. DNA specimens in traces CC approximately 10 ng in mass) can be used for individual discrimination of eukaryotes using the primer in a polymerase chain reaction (PCR).

ANAZ25143 to ANZ25191 represent specifically claimed examples of primers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              eukaryote; restricted identification; DNA sp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Oncorhynchus; restriction primer; short interspersed repeated sequence; eukaryote; restricted polymerase chain reaction fingerprinting; identification; DNA specimen; discrimination; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    repeated sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Restriction primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1999-583348/50.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10-JUL-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; short interspersed repetitive element; SINE; PCR; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human short interspersed repetitive element PCR primer #1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAZ25143 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (NORQ ) NORINSUISANSHO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           868 GGATTACAGGCGTGAGCCAC 887
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                                                                  present
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3; 17pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 98JP-00195692
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                                                                  invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    of eukaryotes
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             for distinguishing individuals with short interspersed of eukaryotes by restricted polymerase chain reaction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SUISANCHO YOSHOKU KENKYUSHOCHO
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1.2e+03;
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RESULT 564
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Best Local (
                                                                                                                                                                           Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAZ25144 standard;
                                                                                                                                                                                                                                                                                  the electrophoresis band pattern, eukaryotic individuals are distinguished. The primer is used for amplifying a eukaryotic decayribonucleic acid (DNA) sequence, pinched between SINE sequences by polymerase chain reaction (PCR) fingerprinting. In particular it may be used individual identification of humans for medical and legal applications and ecological studies. DNA specimens in traces (approximately 10 ng in mass) can be used for individual discrimination of eukaryotes using the primer in a polymerase chain reaction (PCR). ARZ25143 to ARZ25191 represent specifically claimed examples of primers
                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention describes a restriction primer for eukaryotic short interspersed repeated sequences (SINE), which has one or more additional bases that are a mismatch to, or are unrelated to, the 3'-terminal end of the SINE. The annealing temperature of the primer to the DNA sequence is kept higher than the fusion temperature of the primer during polymerase chain reaction (PCR). The PCR fragments obtained are subjected to electrophoresis to obtain a fingerprint. By comparing the polymorphs from electrophoresis to obtain a fingerprint. By comparing the polymorphs from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human short interspersed repetitive element PCR primer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Restriction primer for distinguishing individuals with short interspersed repeated sequence of eukaryotes by restricted polymerase chain reaction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
Homo sapie
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1999-583348/50.
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 ADG70428;
                               ADG70428 standard; DNA;
                                                                                                                                                                                                                                        Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            dentification; DNA specimen; discrimination; ss.
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                                                                                                                                           868 GGATTACAGGCGTGAGCCAC 887
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  6; Page 3; 17pp; Japanese.
                                                                                                             μ.
                                                                                                                                                                                         Similarity
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                                                                                                                                                                                                                                                                        present invention
                                                                                                               GGATTACAGGCGTGAGCCAC 20
                                                                                                                                                                                                                                         BP; 5
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                                                                                                                                                                                                                                         A;
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100.0%; Pred. No.
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                               21 BP
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hes 0;
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RESULT 565
ADG70427
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                                                                                                                                                                                                                                                                                                                                        Creation relates to a novel isolated or recombinant nucleic acid sequence comprising an ANGE, CLLD8 or CLLD7 mRNA, or ANGE-CLD8, ANGE-CLLD8, CLLD7-CLLD8, or ANGE-CLLD8 or CLLD7, CLLD7-CLLD8, or ANGE-CLLD8 or CLLD7 mRNA, or ANGE-CLLD8, ANGE-CLCD8, ANGE-CLLD8 mrd and an activities: antiallergic, antiasthmatic, dermatological, contipyretic, and antiinflammatory. The nucleic acid sequences be used in gene therapy to treat disorders. The nucleic acid sequences care useful for screening agents that inhibit or enhance activity of an ange, CLLD8 or CLLD7 gene. The agent or antibody is useful for treating consecutives, allergic rhinitis or non-atopic asthma. The antibody is useful in an assay detecting or measuring the polypeptide in the sample. The host cell is useful for producing, regulating and analyzing the colypeptide. The splice variant of ANGE, CLLD8, or CLLD7 is useful for CL diagnosing an IgE-mediated disease, atopy, a form of atopic disease or con-atopic asthma, or predicting the severity, or predisposition to a client to the invention.
                                                                                                                                                                                                                                Matches 20;
                                                                                                                                                                                                                                                Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New nucleic acid sequence comprising an ANGE, CLLD8 or CLLD7 mRNA, or their hybrid, useful for screening agents for treating IgE-mediated diseases, e.g. asthma, atopy, hay fever, eczema, atopic dermatitis, or allergic rhinitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21-JUN-2001; 2001GB-00015211.
21-JUN-2001; 2001GB-00015212.
21-JUN-2001; 2001GB-00015213.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ANGE-CLLD8-CLLD7; antiallergic; antiasthmatic; dermatological; antipyretic; antiinflammatory; gene therapy; IgE-mediated disease; REN-34; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       11-MAR-2004 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ANGE; CLLD8; CLLD7; ANGE-CLLD8; ANGE-CLLD7; CLLD7-CLLD8; ANGE-CLLD8-CLLD7; antiallergic; antiasthmatic; dermatolo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REN-34 SNP binding area oligo #2.
                                                                                                                                                                                                                                                                                                Sequence 21 BP; 5 A; 5 C; 9 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 429; 429pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2003-201405/19.
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                 11-MAR-2004
                                                                                                                                                                                                                                                                                                                               oligo relating to the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (ISIS-) ISIS INNOVATIONS LTD
                                                ADG70427;
                                                                               ADG70427 standard; DNA; 21 BP
                                                                                                                                                                                                                                              Local
                                                                                                                                                                                               685 CTCTGCCTCCCGGGTTCAAG 704
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                                                                                                                                                                                                                                              Similarity 100.0%;
                                                                                                                                                              CTCTGCCTCCCGGGTTCAAG 2
                                                                                                                                                                                                                                Conservative
                 (first entry)
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                                                                                                                                                                                                                                                                 2.0%;
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Pred. No.
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REN-34 SNP binding

area oligo

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RESULT 566
ADO11941/c
ID ADO119
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AC ADO119
XC ADO119
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to a novel isolated or recombinant nucleic acid computed an ANGE, CLLD8 or CLLD7 mRNA, or ANGE-CLLD8, ANGE-CLLD8, CLLD7-CLLD8, or ANGE-CLLD8 or CLLD7 mRNA, or ANGE-CLLD8, ANGE-CLLD8 or CLLD7, CLLD7-CLLD8, or ANGE-CLLD8 or CLLD7 hybrid mRNA sequence, its complement, homologue or fragment. The novel nucleic acid sequences have the following activities: antiallergic, antiasthmatic, dermatological, cantipyretic, and antiinflammatory. The nucleic acid sequences are useful for screening agents that inhibit or enhance activity of an CC ANGE, CLLD8 or CLLD7 gene. The agent or antibody is useful for treating CC IgE-mediated diseases, such as asthma, atopy, hay fever, eczema, atopic CC dermatitis, allergic rhinitis or non-atopic asthma. The antibody is useful for producing, regulating and analyzing the colypeptide. The splice variant of ANGE, CLLD8, or CLLD7 is useful for CC diagnosing an IgE-mediated disease, atopy, a form of atopic disease or colion-atopic asthma, or predicting the severity, or predisposition to a colinease. This polynucleotide sequence represents an REN-34 SNP binding colinease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
   ss; primer; simultaneous amplification;
                                                             Single multiplex PCR primer #1313.
                                                                                                                              15-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 429; 429pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               diseases, e.g. asthma, atopy, hay fever, eczema, atopic allergic rhinitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New nucleic acid sequence comprising an ANGE, CLLD8 or CLLD7 mRNA, or their hybrid, useful for screening agents for treating IgE-mediated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-201405/19.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21-JUN-2001; 2001GB-00015211.
21-JUN-2001; 2001GB-00015212.
21-JUN-2001; 2001GB-00015213.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO2003000727-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ANGE; CLLD8; CLLD7; ANGE-CLLD8; ANGE-CLLD7; CLLD7-CLLD8; ANGE-CLLD8-CLLD7; antiallergic; antiasthmatic; dermatoloantipyretic; antiinflammatory; gene therapy; IgE-mediate
                                                                                                                                                                                                                                                          ADO11941 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21-JUN-2002; 2002WO-GB002859.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (ISIS-) ISIS INNOVATIONS LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 685 CTCTGCCTCCCGGGTTCAAG 704
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP; 2 A; 9 C; 5 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                           (first entry)
                                                                                                                                                                                                                                                             DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2.0%; Score 20;
100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cookson
                                                                                                                                                                                                                                                             21
                                                                                                                                                                                                                                                             쁌
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
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                                                                                                                                                                                                                                                                                                                                                                                                                      20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1.2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IgE-mediated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         dermatological;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             dermatitis, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           <u>.</u>.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              읁
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Designing primers for simultaneous amplification of target DNA fragments in a single multiplex polymerase chain reaction, for high throughput multiplex DNA sequence amplification, comprises aligning two primers.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             single multiplex polymerase chain reaction; multifactorial disease;
genetic alteration; pharmacogenetic reaction; genotyping; polymorph
Disclosure; Page 39; 120pp; English.
                                                                                                                                                                                                                                         WPI; 2004-340914/31.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               07-OCT-2002; 2002US-0417009P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              07-OCT-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO2004033649-A2
                                                                                                                                                                                                                                                                                                                                                                                                    (UYNE-) UNIV NEW JERSEY MEDICINE & DENTISTRY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2003WO-US031874.
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The invention relates to a method of designing primers for simultaneous C amplification of target DNA fragments in a single multiplex polymerase CC chain reaction by aligning a first primer and a second primer. The method CC comprises: (a) aligning a first primer and a second primer; and (b) CC selecting the first primer where the first primer at its 3' end does not contain four or more bases that are perfectly matching to the 3' end CC sequence of the first primer or a second primer, the 3' end cC second primer, the first primer at its 3' end does not contain seven or more bases that are perfectly matching CC second primer, the first primer at its 3' end does not contain seven or more bases that are perfectly matching CC first primer or the second primer; and the first primer at its 3' end does not contain eleven or more bases that are perfectly matching to a sequence anywhere of the first primer at its 3' end come not contain eleven or more bases that are perfectly matching except come mismatch to a sequence anywhere of the first primer or the second come mismatch to a sequence anywhere of the first primer or the second come mismatch to a sequence anywhere of the first primer or the second come mismatch to a sequence anywhere of the first primer or the second complification of target DNA fragments in a single multiplex polymerase contain eleven or more bases that are perfectly matching except chain reaction. It is also useful in the identification of multiple genes contain eleven or more bases that are perfectly matching enesic contains the studies in pharmacogenetic reactions, the genotyping genetic contains in a large population, the gene expression invertion. Sequence sequence corresponds to an example of a primer of the 21 BP; 4 A; 6 C; 6 G; 5 T; 0 U; 0 Other; invention.

Ş Matches Query Match Local 545 20; Similarity AGCCTCCCAAGTAGCTGGGA 564 2.0%; llarity 100.0%; Conservative 0, Score 20; Pred. No. Mismatches DB 1; 1.2e+03; 0 Length 21; Indels 0, Gaps

0

RESULT 567
AAZZ5153
ID AAZZ515
XX AAZZ51
XX AAZZ51
XX I3-DEC
DT 13-DEC
XX UBB Human
XX Human AAZ25153; AAZ25153 standard; DNA;

밁

20

AGCCTCCCAAGTAGCTGGGA 1

Human short interspersed repetitive element PCR primer #11

(first entry)

Human; short interspersed repetitive element; SINE; PCR; primer; Oncorhynchus; restriction primer; short interspersed repeated sequence; eukaryote; restricted polymerase chain reaction fingerprinting;

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AAZZ5148
AAZZ5148
AAZ
XX
AC
AAZ
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DT
13-I
XX
Hum
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 568
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Best Local S
Matches 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          bases that are a mismatch to, or are unrelated to, the 3'-terminal end of the SIME. The annealing temperature of the primer to the DNA sequence is kept higher than the fusion temperature of the primer during polymerase chain reaction (PCR). The PCR fragments obtained are subjected to electrophoresis to obtain a fingerprint. By comparing the polymorphs from the electrophoresis band pattern, eukaryotic individuals are distinguished. The primer is used for amplifying a eukaryotic deoxyribonucleic acid (DNA) sequence, pinched between SINE sequences by polymerase chain reaction (PCR) fingerprinting. In particular it may be used individual identification of humans for medical and legal applications and ecological studies. DNA specimens in traces applications and ecological studies. DNA specimens in traces (BCR).

ANZ25143 to ANZ25191 represent specifically claimed examples of primers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Restriction primer for distinguishing individuals with short interspersed repeated sequence of eukaryotes by restricted polymerase chain reaction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         28-JUN-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 6; Page 3; 17pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10-JUL-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       identification; DNA specimen; discrimination; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   repeated
                                                                                                                                                                                                        Oncorhynchus; restriction primer; short interspersed repeated seukaryote; restricted polymerase chain reaction fingerprinting; identification; DNA specimen; discrimination; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
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      28-JUN-1999
                                                                                                                                                                                                                                                                                                   Human; short interspersed repetitive element; SINE; PCR; primer;
                                                                                                                                                                                                                                                                                                                                                           Human short interspersed repetitive element PCR primer #6
                                                                                                                                                                                                                                                                                                                                                                                                                       13-DEC-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAZ25148 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       interspersed repeated
                                                              JP2913035-B1
                                                                                                                                                         Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         868
                                                                                                                      sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GGATTACAGGCGTGAGCCAC 887
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GGATTACAGGCGTGAGCCAC 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP; 5 A; 5 C; 7 G; 5 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              describes a restriction primer for eukaryotic short sequences (SINE), which has one or more additional atch to, or are unrelated to, the 3'-terminal end of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 20;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1; L. 1.2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
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                                                                                                                                                                                                                                                                      sequence;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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RESULT 569
AAZ25154
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       á
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention describes a restriction primer for eukaryotic short conterporated repeated sequences (SINE), which has one or more additional CD bases that are a mismatch to, or are unrelated to, the 3'-terminal end of CD the SINE. The annealing temperature of the primer to the DNA sequence is CD the SINE. The annealing temperature of the primer during polymerase CD chain reaction (PCR). The PCR fragments obtained are subjected to CD cleatrophoresis to obtain a fingerprint. By comparing the polymorphs from CD the electrophoresis band pattern, eukaryotic individuals are CD distinguished. The primer is used for amplifying a eukaryotic construction and ecclosical studies. DNA specimens SINE sequences by CD olymerase chain reaction (PCR) fingerprinting. In particular it may be used individual identification of humans for medical and legal CD (approximately 10 ng in mass) can be used for individual discrimination of eukaryotes using the primer in a polymerase chain reaction (PCR). ANZES143 to AAZES191 represent specifically claimed examples of primers
DR YRA XXX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                        Human; short interspersed repetitive element; SINE; PCR; primer; Oncorhynchus; restriction primer; short interspersed repeated sequence; eukaryote; restricted polymerase chain reaction fingerprinting; identification; DNA specimen; discrimination; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 22 BP; 6 A; 6 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 6; Page 3; 17pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Restriction primer for distinguishing individuals with short interspersed repeated sequence of eukaryotes by restricted polymerase chain reaction
                                                                                                                                                                                                                                                                                                                        Human short interspersed repetitive element PCR primer #12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (NORQ ) NORINSUISANSHO SUISANCHO YOSHOKU KENKYUSHOCHO
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                                                                                                                                                                                                                                                                                                                                                          13-DEC-1999
WPI; 1999-583348/50
                              (NORQ ) NORINSUISANSHO SUISANCHO YOSHOKU KENKYUSHOCHO
                                                                                                                              28-JUN-1999.
                                                                                                                                                             JP2913035-B1
                                                                                                                                                                                             Homo
                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                           AAZ25154 standard; DNA; 22 BP
                                                               10-JUL-1998;
                                                                                               10-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              868 GGATTACAGGCGTGAGCCAC 887
                                                                                                                                                                                           sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GGATTACAGGCGTGAGCCAC 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                            (first entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            98JP-00195692
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            98JP-00195692
                                                                98JP-00195692
                                                                                               98JP-00195692
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 20;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1.2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0,
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RESULT 570
AAZ25150
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention describes a restriction primer for eukaryotic short clinterspersed repeated sequences (SINE), which has one or more additional bases that are a mismatch to, or are unrelated to, the 3'-terminal end of the SINE. The annealing temperature of the primer to the DNA sequence is kept higher than the fusion temperature of the primer during polymerase chain reaction (PCR). The PCR fragments obtained are subjected to electrophoresis to obtain a fingerprint. By comparing the polymorphs from the electrophoresis band pattern, eukaryotic individuals are clistinguished. The primer is used for amplifying a eukaryotic deoxyribonucleic acid (DNA) sequence, pinched between SINE sequences by polymerase chain reaction (PCR) fingerprinting. In particular it may be used individual identification of humans for medical and legal clastions and ecological studies. DNA specimens in traces (approximately 10 ng in mass) can be used for individual discrimination of eukaryotes using the primer in a polymerase chain reaction (PCR). AA225143 to AA225143 to PCR) in the primer in a polymerase chain reaction form the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local S
Matches 20
                                                                                           Restriction primer
                                                                                                                            WPI; 1999-583348/50
                                                                                                                                                                                                                       10-JUL-1998;
                                                                                                                                                                                                                                                        28-JUN-1999
                                                                                                                                                                                                                                                                                    JP2913035-B1
                                                                                                                                                                                                                                                                                                                                      Synthetic.
                                                                                                                                                                                                                                                                                                                                                                               Human; short interspersed repetitive element; SINE; PCR; primer; Oncorhynchus; restriction primer; short interspersed repeated sequence; eukaryote; restricted polymerase chain reaction fingerprinting;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human short
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13-DEC-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAZ25150 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 6; Page 3; 17pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Restriction primer for distinguishing individuals with short interspersed repeated sequence of eukaryotes by restricted polymerase chain reaction
                                                                            repeated
                                                                                                                                                                                         10-JUL-1998;
                                                                                                                                                          (NORQ ) NORINSUISANSHO SUISANCHO YOSHOKU KENKYUSHOCHO
                                                                                                                                                                                                                                                                                                                                                                     identification;
                                                                                                                                                                                                                                                                                                                    sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          868
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20;
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                                                                            sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GGATTACAGGCGTGAGCCAC 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GGATTACAGGCGTGAGCCAC 887
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22
                                                                                                                                                                                                                                                                                                                                                                                                                                                interspersed repetitive element PCR primer #8.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP; 5 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                         98JP-00195692.
                                                                                                                                                                                                                        98JP-00195692
                                                                                                                                                                                                                                                                                                                                                                     DNA
                                                                          r for distinguishing individuals with short in of eukaryotes by restricted polymerase chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       6 C; 7 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ₽₽
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        <u>.</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 20;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 1;
1.2e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
                                                                            interspersed
in reaction
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The present invention describes a restriction primer for eukaryotic short

Claim 6;

Page 3; 17pp; Japanese

fingerprinting.

Claim 6;

Page 3; 17pp; Japanese

Restriction primer

ion primer for distinguishing individuals with short interspersed sequence of eukaryotes by restricted polymerase chain reaction

WPI; 1999-583348/50

(NORQ) NORINSUISANSHO SUISANCHO YOSHOKU KENKYUSHOCHO

repeated

The present invention describes a restriction primer for eukaryotic short interspersed repeated sequences (SINE), which has one or more additional bases that are a mismatch to, or are unrelated to, the 3'-terminal end of the SINE. The annealing temperature of the primer to the DNA sequence is kept higher than the fusion temperature of the primer during polymerase chain reaction (PCR). The PCR fragments obtained are subjected to electrophoresis to obtain a fingerprint. By comparing the polymorphs from the electrophoresis band pattern, eukaryotic individuals are distinguished. The primer is used for amplifying a eukaryotic

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bases that are a mismatch to, or are unrelated to, the 3'-terminal end of the SINE. The annealing temperature of the primer to the DNA sequence is kept higher than the fusion temperature of the primer during polymerase chain reaction (PCR). The PCR fragments obtained are subjected to electrophoresis to obtain a fingerprint. By comparing the polymorphs from the electrophoresis band pattern, eukaryotic individuals are distinguished. The primer is used for amplifying a eukaryotic deoxyribonucleic acid (DNA) sequence, pinched between SINE sequences by polymerase chain reaction (PCR) fingerprinting. In particular it may be used individual identification of humans for medical and legal applications and ecological studies. DNA specimens in traces (approximately 10 ng in mass) can be used for individual discrimination of eukaryotes using the primer in a polymerase chain reaction (PCR).

ARZ25193 to ARZ5191 represent specifically claimed examples of primers
Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                interspersed repeated sequences (SINE), which has one or more additional
                                                                                the present invention
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BP;
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5 C;
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e.
    4 T;
0 U;
    0 Other;
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Matches
                                  Query Match
                          Local
868 GGATTACAGGCGTGAGCCAC
                l Similarity
20; Conser
                 Conservative
                        2.0%;
                 0;
                          Score 20;
Pred. No.
887
                 Mismatches
                                  DB 1;
                          1.2e+03;
                                  Length
                  Indels
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Gaps

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GGATTACAGGCGTGAGCCAC

20

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RESULT 571
AAZ25151
ID AAZ251
                                                                                                                       JP2913035-B1
                                                                                                                                                                                 Synthetic.
                                                                                                                                                                                                                       Oncorhynchus; restriction primer; short interspersed repeated sequence; eukaryote; restricted polymerase chain reaction fingerprinting; identification; DNA specimen; discrimination; ss.
                                                                                                                                                                                                                                                                                                                                                                                                       AAZ25151;
                                                                                                                                                                                                                                                                                                                                                                                                                                            AAZ25151 standard; DNA;
10-JUL-1998;
                                   10-JUL-1998;
                                                                               28-JUN-1999
                                                                                                                                                                                                                                                                                 Human; short
                                                                                                                                                                                                                                                                                                                       Human short interspersed repetitive element PCR primer #9.
                                                                                                                                                                                                                                                                                                                                                                 13-DEC-1999
                                                                                                                                                            domo sapiens.
                                                                                                                                                                                                                                                                                 interspersed repetitive element;
                                                                                                                                                                                                                                                                                                                                                               (first entry
                                       98JP-00195692
98JP-00195692
                                                                                                                                                                                                                                                                                                                                                                                                                                              22
                                                                                                                                                                                                                                                                                                                                                                                                                                              ВP
                                                                                                                                                                                                                                                                                    SINE;
                                                                                                                                                                                                                                                                                 primer;
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Trans and die 12 than

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RESULT 572
AAZZS147
AD AAZZS147
XX AAZZS1
XX AAZZS1
XX AAZZS1
XX I ADEK
DT 13-DEK
DT 13-DEK
DT 13-DEK
XX HUMAN
XX HUMAN
XX HOMON
XX I ONO
XX Synth
OS Synth
OS HOMO
XX I O-JI
XX I OOK

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Best Local S
Matches 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       deoxyribonucleic acid (DNA) sequence, pinches because the may be polymerase chain reaction (PCR) fingerprinting. In particular it may be used individual identification of humans for medical and legal applications and ecological studies. DNA specimens in traces applications and ecological studies bused for individual discrimination (Androximately 10 ng in mass) can be used for individual discrimination (PCR).
                                                                                                                                                                              The present invention describes a restriction primer for eukaryotic short interspersed repeated sequences (SIME), which has one or more additional bases that are a mismatch to, or are unrelated to, the 3'-terminal end of the SINE. The annealing temperature of the primer to the DNA sequence is kept higher than the fusion temperature of the primer during polymerase chain reaction (PCR). The PCR fragments obtained are subjected to electrophoresis to obtain a fingerprint. By comparing the polymorphs from the electrophoresis band pattern, eukaryotic individuals are distinguished. The primer is used for amplifying a eukaryotic deoxyribonucleic acid (DNA) sequence, pinched between SIME sequences by polymerase chain reaction (PCR) fingerprinting. In particular it may be used individual identification of humans for medical and legal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (approximately 10 ng in mass) can be used for individual discrimination of eukaryotes using the primer in a polymerase chain reaction (PCR).

AAZ25143 to AAZ25191 represent specifically claimed examples of primers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human short interspersed repetitive element PCR primer #5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; short interspersed repetitive element; SINE; PCR; primer; Oncorhynchus; restriction primer; short interspersed repeated sequence; eukaryote; restricted polymerase chain reaction fingerprinting;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                13-DEC-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Restriction primer for distinguishing individuals with short interspersed repeated sequence of eukaryotes by restricted polymerase chain reaction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    JP2913035-B1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 6; Page 3; 17pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1999-583348/50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (NORQ ) NORINSUISANSHO SUISANCHO YOSHOKU KENKYUSHOCHO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10-JUL-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              identification;
                                     (approximately 10 ng in mass) can be used for individual discrimination of eukaryotes using the primer in a polymerase chain reaction (PCR). AAZ25143 to AAZ25191 represent specifically claimed examples of primers
                                                                                                                                                    used individual identification of humans for medical and legations and ecological studies. DNA specimens in traces
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        the present invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       98JP-00195692
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA specimen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  discrimination;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  88.
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RESULT 573
ABL55369
ID ABL553
XX CANCAL
DT 23 -JUL
XX CANCAL
PN CN1331
XX CN1331
XX CN1331
XX ABL553
XX AB
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Best Local Similarity
Thes 20; Conserv
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밁
                                             S
                                                                                                  Matches
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Best Local
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                                                                                                                                                                                                                                                                                             The invention relates to human leucine zipper protein 11.99 (AAM49285) and to nucleic acids encoding it (ABL55389). The protein has a molecular weight of 12 kD. The invention also relates to a method for the recombinant production of the protein, an antagonist of the protein, and the use of the protein, gene and antagonist in therapeutic applications. Leucine zipper protein 11.99 can be used in the treatment of a variety of diseases such as embryonic development disorders and tumours. Sequences ABL55399-ABL55370 represent reverse transcription-PCR (RT-CR) primers and the control of the protein 11.99 can be used in the treatment of a variety of the protein as embryonic development disorders and tumours. Sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; leucine zipper protein 11.99; recombinant cancer; embryonic development disorder; cytostat:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       30-JUN-2000; 2000CN-00116898
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-JUN-2000; 2000CN-00116898
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16-JAN-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CN1331194-A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            23 -JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABL55369 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Polypeptide-human leucine zipper protein 11.99 and polynucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-292862/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       мао Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      reverse
                                                                                                                                                                                                  Sequence 24 BP; 4 A; 5 C; 10 G; 5 T; 0 U; 0 Other
                                                                                                                                                                                                                                               zipper protein 11.99
                                                                                                                                                                                                                                                                             used in an exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (BODE-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           868 GGATTACAGGCGTGAGCCAC
                                               943
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            leucine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 μ.
                                                                                                  20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    embryonic development disorder; cytost
transcription-PCR; RT-PCR; primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Xie
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BODE GENE
                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GGATTACAGGCGTGAGCCAC
                                                  CCCAGGCTGGAGTGCAATGG 962
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 6 A;
                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      zipper
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19 (Disclosure); 35pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DEV CO LTD SHANGHAI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   5 C; 7 G; 4 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%;
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                                                                                                                             100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                     0
                                                                                                                          Score 20;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            11.99 RT-PCR primer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 887
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                                                                                                     Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            combinant production; tumour;
cytostatic; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1;
                                                                                                                                                     DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1.2e+03
                                                                                                                             1.3e+03;
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                                                                                                                                                  Length 24;
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                                                                                                          Indels
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RESULT 574 ADC56863/c ID ADC568 XX

ADC56863 standard; DNA;

24

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RESULT 575
AAH77599
ID AAH775
XX AAH775
XX AC AAH775
XX Human
XX Human;
KW human;
KW human;
KW inflam
XX Homo 8
XX CO1298
XX C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This invention relates to a novel protein, human protein 8-91, and the DNA sequence encoding it. The protein of the invention may be useful the treatment of diseases such as diabetes and cancer. The present sequence is that of an RT-PCR primer which was used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18-APR-2001; 2001CN-00112644.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            27-NOV-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     reverse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             human;
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                                                                                                                   24-NOV-1999;
                                                                                                                                                                06-JUN-2001.
                                                                                                                                                                                                              CN1298002-A.
                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                  human immunodeficiency virus; HIV; infection; immunological disease; inflammatory disease; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAH77599
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      exemplification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Polypeptide-human ribosomal protein -8.91
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18-APR-2001; 2001CN-00112644.
                          (SHAN-) SHANGHAI BORONG GENE DEV
                                                                       24-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                  Human; dihydropyrrole-5-carboxylate reductase 30; cancer; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                             Human dihydropyrrole-5-carboxylate reductase
                                                                                                                                                                                                                                                                                                                                                                                                                                             22-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAH77599 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    868
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e transcription PCR; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      primer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GGATTACAGGCGTGAGCCAC 887
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP; 4 A; 8 C; 6 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                             (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
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                                                                       99CN-00124090
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ID NO 3;
                                                                                                                   99CN-00124090
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                             entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ₽₽.
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    and
                                                                                                                                                                                                                                                                                                                                                                                               30 PCR primer
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RESULT 576
AAF74080
ID AAF740
XX AAF740
XX Solute
AC AAF740
XX Solute
KW Genoty
XX Homo 8
XX Homo 8
XX Homo 8
XX Homo 8
XX Homo 9
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ID AAF740
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Best Local S
Matches 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    genotyping; allele
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Solute carrier family 6 neurotransmiter transporter; sectonin genotyping; allele specific oligonucleotide; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 25 BP; 7 A; 1 C; 5 G; 12 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                   New isolated polynucleotide comprising a polymorphic variant for the solute carrier family 6 neurotransmitter transporter, serotonin member 4 gene for identifying drugs for treating disorders related to expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       29-JUL-1999;
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                                                                                                                                                                                                                                         Example 1; Page
                                                                                                                                                                                                                                                                                             of the protein.
                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-123317/13
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                                                                                                                                                                                                                                         33; 152pp;
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                                                                                                                                                                                                                                         English.
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The present invention relates to a polymorphic variant of a reference sequence for the solute carrier family 6 neurotransmitter transporter, serotonin member 4 (SLC6A4) gene or a fragment of it or a sequence complementary to the first sequence. The invention is used in producing recombinant organism that can be used to express SLC6A4 for protein structure analysis and binding studies. A composition comprising a

structure analysis and binding studies. A congenetyping oligonucleotide is used to detect

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                                                                                        Query Match
Best Local Similarity
Matches 21; Conserv
                                                                                                                                                                                                                                      The invention relates to a novel method for extracting intact cytoplasmic biomolecules from cells. The method of the invention is useful for extracting or acquiring cytoplasmic biomolecules such as proteins or nucleic acids which include cytoplasmic RNA, nuclear and mitochondrial RNA, nuclear and mitochondrial DNA, cytoplasmic mRNA, or their combinations from cells. The method is useful in cancer screening, selecting and monitoring for chemotherapy treatment or cancer recurrence. This type of cell analysis is useful in cancer diagnostics. The method is useful in profiling cells isolated from tissues or body fluids and serves as an adjunct to clinical diagnosis of diverse carcinomas including early stage detection and classification of circulating tumour cells. The present sequence is used in the exemplification of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Extracting intact cytoplasmic biomolecules e.g. proteins, nucleic from cells, by treating sample comprising cells containing target with permeabilizing agents to release biomolecules and recovering
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     26-OCT-2001; 2001US-0330669P 04-APR-2002; 2002US-0369945P
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                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 10; Page 59; 119pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CC in a test subject, by obtaining a sample having a mixed cell population CC in a test subject, by obtaining a sample having a mixed cell population CC suspected of containing cancer cells of epithelial origin, mixing the CC sample with immunomagnetic particles which bind specifically to the CC cancer cells, subjecting the mixture to produce a separated cell CC cancer cells. The method is useful for diagnosing the severity of a CC cancer cells. The method is useful for diagnosing the severity of a CC presence of circulating cancer cells. The test subject is for assessment of a CC cancer cells. The test subject is for assessment of a CC cancer cells. The test subject is subject response to cancer CC cancer cells. The test subject has been diagnosed with a cancer selected from prostate cancer, breast cancer, colon cancer apudoma, choristoma, CC braichoma, malignant carcinoid syndrome, carcinoid heart disease, and CC carcinoma e.g. Walker, basal cell, basosquamous, Brown-Pearce, ductal, CC chis sequence represents a primer used to amplify a specific gene cDNA corcer in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                 Matches
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       29-JAN-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28-OCT-2002; 2002WO-US034570
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 10; Page 59; 105pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2003-421425/39.
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                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C; 6
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                                                                                                                                                                                                                                                                                                                                                                                 Score 19.8; D. Pred. No. 1.2e. 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        G; 5 T;
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ARESULT 580
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DE PCR pr
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                                                                                                                                                                                                                                                                                                                                         containing-1 (SORBS1) gene. The method or determining the containing-1 (SORBS1) gene. The method comprises determining the containing-1 (SORBS1) gene or more positions chosen from 220; 249; -7 with respect to exon 5; -25 with respect to exon 10; +69 with respect to exon 11; +33 with respect to exon 12; and 2337. The invention also discloses primer sequence by amplification and sequencing of the gene. The method is useful for associating one or more SORBS1 SNPs with an insulin disorder e.g. type 2 diabetes, obesity, hypertension, atherosclerosis or metabolic syndrome. The presence or absence of the SNP may be useful in determining whether an individual is at increased or decreased risk for an insulin disorder. The SNPs were identified by screening all of the exons, and 50-150 base pairs of the flanking regions of the introns of the SNP in the human SORBS1 gene. The present sequence represents a sequencing primer used to sorder the human SORBS1 gene.
                                                                                                                                                                                                                                                   Matches
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Best Local (
 PCR primer for human DNA marker clone G212.
                                   23-DEC-1999
                                                                   AAZ27796;
                                                                                                  AAZ27796
                                                                                                                                                                                                                                                                                                                Sequence 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 1; Page 8; 18pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         atherosclerosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Detecting at least one single nucleotide polymorphism in a human and SH3-domain-containing-1 (SORBS1) gene, useful in diagnosing disorders like type 2 diabetes, obesity, hypertension and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (HSIA/)
(TAIT/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sorbin and SH3-domain-containing-1 gene; SORBS1; sequence determination; insulin disorder; type 2 diabetes; obesity; hypertension; atherosclerosis; metabolic syndrome; sequencing; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Single nucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hsiung CA,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          present invention relates to a method of detecting at least one gle nucleotide polymorphism (SNP) in a human sorbin and SH3-domain-
                                                                                                                                                                                                                 870
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SORBS1
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                                                                                                                                                                                                                                                                Similarity
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                                                                                                standard;
                                                                                                                                                                                 ATTACAGGCATGAGCCACCACAC 23
                                                                                                                                                                                                               ATTACAGGCGTGAGCCACCACGC 892
                                                                                                                                                                                                                                                 Conservative
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                                 (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   polymorphism;
                                                                                                  DNA;
                                 entry)
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                                                                                                                                                                                                                                                                                                                8 C; 4 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                               Score 19.8;
Pred. No. 1
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AAA46454;

04-SEP-2000

(first entry)

RESULT 581 AAA46454/c

AAA46454 standard; DNA; 24

ВP

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Query Match
Best Local Similarity
Matches 21; Conserv

Conservative

2.0%;

Score 19.8; I Pred. No. 1.3e 0; Mismatches

1.3e+03

DB 1;

Length

24;

0,

Gaps

0

638 23

TGTCACCCAGGCTGGAGTGCAGT 660

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CC hybridization selection, and comprises: (a) providing several DNA CC fragments, at least one of which contains an ITR sequence, a region of the DNA fragment which contains at least one repeat unit consisting of a CC sequence of five, six or seven bases repeated in tandem at least two contains; the description of the providing a stationary support having at least one cC cligonucleotide associated with it, where the oligonucleotide includes a cC sequence of nucleotides which is complementary to a portion of the ITR cc sequence, and (c) combining the DNA fragments with the support under CC conditions where the DNA fragments including the DNA fragment containing the ITR sequence hybridize to the support. The method is particularly used to isolate DNA containing pentanucleotide tandem repeat sequences as cell as to detect target ITR DNA sequences having a low incidence of CC stutter artifacts (no more than 2.4%). The method is useful in DNA CC profiling for linkage analysis, criminal justice, paternity testing and CC the lineage of horses, dogs and other prize animals. The invention CC profiling. The method can detect polymorphisms with a low incidence of Stutter artifacts, which has previously been a problem in interpreting CC allelic content of loci. The development of markers based on larger repeat units, enables easier separation of the fragments of more loci or content of loci. The development of markers based on larger calculations and content of loci. The development of markers based on larger calculations are content of loci. The development of markers based on larger calculations are calculated and content sequences as calculated and content sequences as calculated and content sequences of the cross celling and content content of loci. The development of markers based on larger calculated and content content
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Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         method of the containing an
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Isolating DNA containing intermediate tandem repeat sequences, useful in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (PROM-) PROMEGA CORP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       sequence
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BP;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     is a PCR primer for a human DNA marker clone used invention. The method is for isolating a fragment intermediate tandem repeat (ITR) sequence using
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       98US-00018584.
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    0 Other
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RESULT 582
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                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local S
Matches 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 are arranged in the same transcriptional orientation. The nucleic acid is used to specifically express the nucleic acid of interest in vivo, particularly in the nervous system and especially expression of tyrosine hydroxylase for treatment of neurodegeneration (Parkinson's disease), nervous system injury and retinal degeneration. More generally, it can be used to express a very wide range of therapeutic products, e.g. enzymes, blood factors, cytokines, tumour suppressors, antibodies etc., for (immuno) therapy of infections, cancer, autoimmune diseases, restenosis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The specification describes a nucleic acid which comprises a region (R1) encoding the transactivator (tTA) of the tetracycline-regulated system, controlled by a moderate promoter; and a region (R2) comprising a nucleic acid of interest under control of a promoter sensitive to tTA. R1 and R2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example; Page 24; 51pp; French
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New nucleic acid for regulating gene expression, particularly of tyrosine hydroxylase for treatment of Parkinson's disease, the gene and tetracycline transactivator.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         03-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Transactivator; tetracycline-regulated system; promoter; nervous tyrosine hydroxylase; neurodegeneration; Parkinson's disease; nervous system injury; retinal degeneration; probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Oligonucleotide
                                                       Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (AVET ) AVENTIS PHARMA SA
                                   SNPE; genotyping; agammaglobulinaemia; dia
Lesch-Nyhan syndrome; muscular dystrophy;
polycystic kidney disease; osteogenesis in
                                                                                                                                                             14-AUG-2001
                                                                                                                                                                                                                                 AAH39521
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    sequences and ribozymes. The present sequence represents identify human cells, in the course of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      genetic diseases etc., also antigens for vaccination or antisense
                                                                                                                            specific upper
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2000-387422/33
                                                                                                                                                                                                                                                                                                                                                        661 GGCGCAATCTTGGCTCACTGCAA 683
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                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP; 4 A; 7 C; 8 G; 4 T; 0 U; 1 Other;
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                                                                                                                                                               (first entry)
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99US-0122600P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              probe used to detect human cells.
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                                                                                                                                                                                                                               DNA;
 porphyria; rheumaco
nsic investigation;
                                                                                                                            PCR primer SEQ ID 2317.
                                                                                                                                                                                                                                                                                                                                                                                                             2.0%;
                                                                                                                                                                                                                                 24
                                                                                                                                                                                                                                                                                                                                                                                               <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                               Pred. No. 1.30
); Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                              Score 19.8;
                    ceogenesis imperfecta; autoimmune disease;
rheumatoid arthritis; multiple sclerosis;
                                                                                                                                                                                                                                                                                                                          N
   paternity
                                                                                                                                                                                                                                                                                                                                                                                                                 .3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                              Length 24;
                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         a probe
   PCR primer; ss
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Crimer extension (SNPE) primers, and the sequences of regions flanking concludes kits for determining the presence or absence of a SNP, using the coligonucleotides of the invention. The PCR primers are used to amplify a CC SNP flanking sequence, the SNPE primer is used as a genotyping primer. CC SNP flanking sequence, the SNPE primer is used as a genotyping primer. CC performing a single-nucleotide primer extension reaction. The CC performing a single-nucleotide primer extension reaction. The CC identity of a SNP and for genotyping nucleic acid sample by CC genotity of a SNP and for genotyping nucleic acid samples, for e.g. to CC identity of a SNP and for genotyping nucleic acid samples, for e.g. to CC assess by association analysis the genotype of an individual or group of CC individuals, having a pathological phenotypic trait suspected of being CC caused by one or more SNPs. Phenotypic traits include diseases e.g. CC osteogenesis imperfecta and acute intermittent porphyria. Phenotypic craites also include symptoms of or susceptibility to multifactorial CC diseases, including, rheumatoid arthritis, multiple sclerosis, inflammation, cancer, nervous system diseases and infection by pathogenic microorganism. The method is also useful in forensic investigations and contentive analysis are received.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequences AAH37205 -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1; Page 61; 83pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         acid sample.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           absence or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2001-290930/30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            13-OCT-2000; 2000WO-US028436
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO200129262-A2
                                     paternity analysis. for a human SNP cont
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New genotyping oligonucleotide, useful for detecting the presence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (ORCH-) ORCHID BIOSCIENCES INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               identity of single polynucleotide polymorphism
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        99US-0160096P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ļ,
                                       containing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pohl M;
                                                       The present sequence represents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAH40944 represent PCR primers, single nucleotide
                                     DNA sequence
                                                             þ
                                                             PCR
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                                                             primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 a nucleic
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Sequence 24 BP; 6 A; 7 C; 3 G; 8 T; 0 U; 0 Other;

Matches Query Match Local 21; Similarity Conservative 91.3%; 2.0%; 0; Mismatches Pred. Score 19.8; NO. 1.3e+03 DB 1; Length 24; Indels 0; Gaps 0

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밁
           577 ACCACTACACCTGGCTAATTTTT 599
ACCACTACGCCTGACTAATTTTT
 23
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AAH4468

AAH44468 standard; DNA;

24 ВP S

Enolpyruvate phosphate-dependent glycophosphotransferase 9 primer 1.

25-OCT-2001

(first entry)

cytostatic; antiinflammatory; haemostatic; immunomodulatory;
diagnosis; malignant neoplasm; haemopathy; HIV infection;
immunological disease; inflammation; PCR primer; ss. Human; enolpyruvate phosphate-dependent glycophosphotransferase anti-HIV;

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RESULT 584
AAI64654/c
ID AAI646
XX AAI646
XX AAI646
XX Human
XX Human;
KW Human;
KW antiin
KW human
XX Homo 8
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XX MAO Y,
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention describes the human enolpyruvate phosphate-dependent glycophosphotransferase 9 protein (1). (1) has cytostatic, antiinflammatory, haemostatic, immunomodulatory and anti-HIV activities. (I) and the polynucleotide encoding it are applicable in the diagnosis and treatment of malignant neoplasm, haemostatly, HIV infection, immunological diseases and various inflammations. The present sequence represents a PCR primer for human enolpyruvate phosphate-dependent glycophosphotransferase 9, which is used in an example from the present
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                                                                                                                    04-OCT-2001.
                                                                                                                                            WO200172971-A1
                                                                                                                                                                                              Human; RNA helicase 10; cytostatic; virucidal; immunomodulatory; antiinflammatory; haemostatic; malignant tumour; HIV; infection; human immunodeficiency virus; immunological disease; PCR primer;
                                                                                                                                                                                                                                                  Human RNA helicase
                                                                                                                                                                                                                                                                            04-DEC-2001
                                                                                                                                                                                                                                                                                                                              AAI64654 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 3; Page 18; 35pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 neoplasm, hemopathy, HIV infection, immunological diseases and various inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Enolpyruvate phosphate-dependent glycophosphotransferase 9 and encoded polynucleotide, applicable in diagnosis and treatment of malignant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-432875/46
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                                                                  27-MAR-2000; 2000CN-00115186
                                                                                          26-MAR-2001; 2001WO-CN000435
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                                         (SHAN-)
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                                        SHANGHAI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SHANGHAI BIO DOOR GENE TECHNOLOGY LTD
                                                                                                                                                                                                                                                                                                                                                                                                                  GGAATCTCACTCTGTTACCCAGG
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
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                                        BIOWINDOW GENE
                                                                                                                                                                                                                                                10 PCR primer 2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 19.8;
Pred. No. 1.
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                                                                                                                                                                                               PCR primer;
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ABL41338
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                            The invention relates to human TFIID subunit p30beta protein 12.54, the polynucleotide encoding this polypeptide and DNA recombinant processes to produce the polypeptide. The present invention also discloses the method of applying the polypeptide in treating various diseases, such as protein metabolism dysfunction, various tumours, inflammations and immunological diseases, haemopathy and HIV infection. The present invention also discloses the antagonist for resisting the polypeptide and its treatment effect. The present invention also discloses the application of the polypurcleotide for encoding human TFIID subunit p30beta protein 12.54.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; TFIID subunit p30beta protein 12.54; tumour; protein metabolism dysfunction; immunological diseas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The polypeptide and encoded polynucleotide are applicable in diagnosis and treatment of malignant tumour, haemopathy, HIV infection, immunological diseases and various inflammations. The present sequence that of a human RNA helicase 10 PCR primer, useful to the invention
                                                                                                                                                                                                                                                                                                                                                                                                  New polypeptide-human TFIID subunit p 30 polynucleotide encoding the polypeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mao
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               05-JUN-2000; 2000CN-00116325
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      immunodeficiency virus infection, immunological diseases and inflammation.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-597114/67.
                                                                                                                                                                                                                                                                                                                                          Example 2; Page 19 (Disclosure); 35pp; Chinese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2002-206968/27.
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Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SHANGHAI.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     disease; haemopathy; HIV;
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The present sequence is that of a PCR primer, useful in examples of the

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RESULT 587
AAD38977
ID AAD389
XX
AC AAD389
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AC AAD389
XX
XX
XX
ZY
ZY
Z3-SEF
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AAI66361
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Best Local S
Matches 21
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Best Local Similarity
Matches 21; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; phosphatidylinositol-3 kinase 35; PTDINS-3 kinase 35; cancer; haemopathy; development disorder; HIV infection; immunological disease; inflammation; gene therapy; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human phosphatidylinositol-3 kinase 35 cDNA PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAI66361 standard; DNA;
                                                                                                                                                                                                                                     The present invention provides the protein and coding sequences of human phosphatidylinositol-3 (PTDINS-3) kinase 35. The sequences can be used in the treatment of cancer, haemopathy, HTV infection, development disorders, immunological diseases and inflammation. The present sequence
                                                                                                                                                                                                                                                                                                                   New human phosphatidylinositol-3 (PTDINS3) kinase 35 for diagnosing treating malignant tumor, hemopathy, human immunodeficiency virus infection, immunological diseases and various inflammations.
                                                                                                                                                                                                                                                                                                                                                                                                                                     17-MAR-2000; 2000CN-00114973.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO2001,75014-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       23-JAN-2002
                                                                                                                                                                                                      Sequence 24
                                                                                                                                                                                                                                                                                                                                                                  WPI; 2002-025836/03.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         16-MAR-2001; 2001WO-CN000328
             23-SEP-2002
                                  AAD38977;
                                                                                                                                                                                                                                                                                               Example 2; Page 12; 34pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                             (BIOW-)
                                                       AAD38977
                                                                                                                                                                                                                            a PCR
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                                                                                                                                                         l Similarity
21; Conserv
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                                                                                                                              TTTTTATTTTATTTTTTTAAGA 449
                                                                                                                                                                                                                            primer for the coding sequence of the invention
                                                      standard;
                                                                                                               TTTTTTTTTTTTTTTAAGA 24
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ilarity 91.3%;
Conservative
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Pred. No. 1
                                                                                                                                                                      Score 19.8;
Pred. No. 1.
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                                                                                                                                                           Mismatches
                                                                                                                                                                      8; DB 1;
1.3e+03;
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RESULT 588
ABL41577
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention relates to dipeptidyl peptidase (DPP) proteins and polynucleotides encoding such proteins. The DPP peptides are useful for screening inhibitors of DPP catalytic activity. The inhibitors are useful for treating neoplasia, type II diabetes, cirrhosis, autoimmunity, graft rejection and HIV (human immuno deficiency virus) infection. The present DNA sequence is a reverse transcription (RT)-PCR primer which is used for amplifying human GDD DNA. This sequence is used in the exemplification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New dipeptidyl peptidase (DPP) peptides, useful for screening inhibitors of DPP catalytic activity, which may be employed to treat e.g. neoplasia. type II diabetes, cirrhosis, autoimmunity, graft rejection and HIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human GDD DNA amplifying reverse RT-PCR primer, GDDpr-4r.
                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 24 BP; 2 A; 7 C; 8 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  02-MAY-2002.
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                                                                                                                                                                                                           01-JUL-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        29-OCT-2001; 2001WO-AU001388.
                        31-MAY-2000;
                                                  31-MAY-2000; 2000CN-00116276
                                                                                                    CN1325869-A.
                                                                                                                              Homo sapiens
                                                                                                                                                       Zinc finger;
                                                                                                                                                                                 Primer #2 relating to human zinc
                                                                                                                                                                                                                                      ABL41577;
                                                                                                                                                                                                                                                                ABL41577
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 the invention
(BODE-) BODE GENE DEV CO LTD SHANGHAI
                                                                           12-DEC-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (UNSY ) UNIV SYDNEY
                                                                                                                                                                                                                                                                                                                                              197 CCATGTTGGTCAGGCTGGTCTCG 219
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                                                                                                                                                                                                                                                                standard;
                                                                                                                                                                                                                                                                                                                                 CCATGTTGGCCAGGCTGGTCTTG
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                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                        2000CN-00116276
                                                                                                                                                        zinc finger protein 27; human; cancer; HIV; PCR; primer;
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                                                                                                                                                                                                                                                                DNA;
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Pred. No. 1
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                                                                                                                                                                                   finger protein
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RESULT 589
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Best Local S
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                                                                                Sequence
                                                                                                  The present invention relates to human RNA polymerase I-40 kDa subuse.68 (see ABP59130). The protein can be used for treating diseases as cancer and HIV infection. The present sequence is a PCR primer, was used in an example from the invention
                                                                                                                                                       Example 2; Page 17 (Disclosure); 32pp; Chinese.
                                                                                                                                                                                     Polypeptide-human RNA polymerase I-40 kDa subunit 9.68 and polynuceotide
                                                                                                                                                                                                                                                                        05-JAN-2001;
                                                                                                                                                                                                                                                                                            05-JAN-2001; 2001CN-00105029.
                                                                                                                                                                                                                                                                                                                  14-AUG-2002.
                                                                                                                                                                                                                                                                                                                                     CN1363655-A.
                                                                                                                                                                                                                                                                                                                                                                               Human; RNA polymerase I-40 kDa subunit 9.68; cancer; cytostatic;
HIV infection; anti-HIV; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                              Human RNA polymerase I-40 kDa subunit 9.68 PCR primer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Polypeptide-zinc finger protein 27 and polynucleotide for coding
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                                                                                                                                                                                                                                                   (BODE-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            s invention relates to a novel polypeptide-zinc finger protein 27 and application of the polypeptide in treating diseases such as cancer HIV infection. The present sequence represents a primer relating to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ķ
                                                                                                                                                                           coding
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                   219
                                                                                                                                                                                                                                                                                                                                                          sapiens.
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                                                 Similarity
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GAACTCCTGACCTCAGGTGATCC 23
            GAACTCCCGACCTCAGATGATCC 241
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ACCACGCCTGGCTAACTTTTTTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ACCACGCCTAGCTAATTTTTTTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              24 BP; 4 A; 7 C; 5 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  finger protein 27 encoding sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Υ,
                                                                               BP; 6 A; 8 C; 5 G; 5 T; 0 U; 0 Other;
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                                        Conservative
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                                                2.0%;
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Pred. No. 1.3e
0; Mismatches
                                                 Score 19.8;
Pred. No. 1.
                                        Mismatches
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2;
                                                .3e+03;
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RESULT 591
ADO11357
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ADL09135
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                            The invention relates to a new human protein phosphatase 2A alpha subunit 22. 66 polypeptide, the polymucleotide encoding it, and the process for preparing this polypeptide by DNA recombination technique. Also described is the application of the said polypeptide in treating diseases such as osteoma and leukaemia; the antagonist against this polypeptide and its therapeutic action; and the application of the said polypucleotide encoding this novel polypeptide. The present sequence represents a reverse transcriptase (RT)-PCR primer used to isolate cDNA encoding human protein phosphatase 2A alpha subunit 22. 66.
                                  single multiplex polymerase chain reaction; multifactorial genetic alteration, pharmacogenetic reaction, genotyping; p
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Polypeptide-human protein phosphatase 2A alpha subunit 22.66 polynucleotide for coding it.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mao
Synthetic
                                                            ss; primer;
                                                                                    Single multiplex PCR
                                                                                                                                                            ADO11357
                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example
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leukaemia; RT-PCR; reverse transcriptase; primer.
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                                                                                                            15-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (UYFU-) UNIV FUDAN
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                       expression
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2; Page 18; 32pp; Chinese
                                                                                                                                                             standard; DNA;
                                                                                                                                                                                                                                                 GGCTCAAGCGATTCTCCTGTCTC 1019
                                                                                                                                                                                                                                                                                                                        24 BP; 4 A; 8 C; 5 G; 7 T; 0 U; 0 Other;
                                                            simultaneous amplification;
                                                                                                                                                                                                                                                                        Conservative
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                                                                                                            (first
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                                                                                                            entry)
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                                                                                    primer
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                                                                                     #729.
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Pred. No. 1
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                                                                                                                                                                                                                                                                                              DB 1;
                                   genotyping; polymorphism;
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                                                                                                                                                                                                                                                                                              Length 24;
                                                                                                                                                                                                                                                                          Indels
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RESULT 592
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Matches
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           more bases that are perfectly matching to a sequence anywhere of the first primer or the second primer, and the first primer at its 3' end does not contain eleven or more bases that are perfectly matching except one mismatch to a sequence anywhere of the first primer or the second primer. The method is useful for designing primers for simultaneous amplification of target DNA fragments in a single multiplex polymerase amplification in the sales useful in the identification of multiple genes related to multifactorial diseases, the genome-scale detection of genetic alterations, the studies in pharmacogenetic reactions, the genotyping genetic polymorphisms in a large population, the gene expression profiling in various samples and high throughput genotyping technologies. This sequence corresponds to an example of a primer of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           amplification of target DNA fragments in a single multiplex polymerase chain reaction by aligning a first primer and a second primer. The method comprises: (a) aligning a first primer and a second primer; and (b) selecting the first primer where the first primer at its 3' end does not contain four or more bases that are perfectly matching to the 3' end sequence of the first primer or a second primer; the first primer at its 3' end does not contain seven or more bases that are perfectly matching except one mismatch to the 3' end sequence of the first primer or the second primer; the first primer at its 3' end does not contain six or second primer; the first primer at its 3' end does not contain six or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Designing primers for simultaneous amplification of target DNA fragments in a single multiplex polymerase chain reaction, for high throughput multiplex DNA sequence amplification, comprises aligning two primers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Li H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   07-OCT-2002; 2002US-0417009P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to a method of designing primers for simultaneous amplification of target DNA fragments in a single multiplex polymerase and a second primer. The method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (UYNE-) UNIV NEW JERSEY MEDICINE & DENTISTRY
                                                                                                                                                                                                           19-FEB-2001
                                                                                                                                                                                                                                                                                     AAC61385
                    WO200058470-A1
                                                                                                              Androgen-regulated gene; prostate specific gene; PCGEM1; prostate cancer; prostate cancer gene expression marker 1; prostate related disease;
                                                                                                                                                                   PCR primer for androgen-related, prostate-specific gene PCGEM1.
                                                        Homo sapiens
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prostatic hype
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                                                                                                                                                                                                                                                                                     standard;
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                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                     DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 19.8; DB 1;
Pred. No. 1.3e+03;
                                                                                            PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
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RESULT 593
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                                                                                                                                                                                                                                                                                                                                          Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis; inflammation; forensic investigation; paternity analysis; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel androgen-regulated prostate specific gene, prostate cancer expression marker, useful for detecting, diagnosing, preventing, treating prostate cancer and other prostate related diseases.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SNP specific lower PCR primer SEQ ID 1406.
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MOUL J W.
SRIVASTAVA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA;
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15-OCT-1999;

13-OCT-2000; 2000WO-US028436.

26-APR-2001

WO200129262-A2

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RESULT 594
AAH38522
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Best Local
                                                                                                                                                                                                                                                                                                    Single nucleotide polymorphism; war; wanger consisting diabetes insipidus; cancer; SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis; section in the scheme of the scheme 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide primer extension (SNPE) primers, and the sequences of regions flanking sites of single nucleotide polymorphisms SNPs. The present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 21
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15-OCT-1999;
                                                       13-OCT-2000; 2000WO-US028436.
                                                                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SNP specific lower PCR primer SEQ ID
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                                                                                                                         26-APR-2001
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99US-0160096P.
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95.2%;
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Pred. No. 1.
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identity of a SNP and for genotyping nucleic acid samples, for e.g. to assess by association analysis the genotype of an individual or group of individuals, having a pathological phenotypic trait suspected of being caused by one or more SNPs. Phenotypic traits include diseases e.g. agammaglobulinaemia, diabetes insipidus, Lesch-Nyhan syndrome, muscular dystrophy, familial hypercholesterolaemia, polycystic kidney disease, osteogenesis imperfects and acute intermittent porphyria. Phenotypic traits also include symptoms of or susceptibility to multifactorial disease of which a component is or may be genetic such as autoimmune diseases, including, rheumatoid arthritis, multiple sclerosis,
                                                                                                                                                                                                                                                                                                                                                                                                                          Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide primer extension (SNPE) primers, and the sequences of regions flanking sites of single nucleotide polymorphisms SNPs. The present invention includes kits for determining the presence or absence of a SNP, using the oligonucleotides of the invention. The PCR primers are used to amplify a SNP flanking sequence, the SNPE primer is used as a genotyping primer. The oligonucleotides are useful for genotyping a nucleic acid sample by performing a single-nucleotide primer extension reaction. The oligonucleotides are useful for determining the presence, absence or oligonucleotides are useful for determining the presence, absence or
                                                                        inflammation, cancer, nervous system diseases and infection by pathogenic microorganism. The method is also useful in forensic investigations and paternity analysis. The present sequence represents a PCR primer specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nucleic acid sample.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 1; Page 56; 83pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Picoult-Newburg L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ORCH-) ORCHID BIOSCIBNCES INC
                                                     human SNP
                                                  containing
                                                  DNA sequence
                                                                                                                                                                                                                                                                                                                                                                             of.
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Query Match Best Local S Matches 20 Sequence l Similarity 20; Conserv 21 BP; 6 Conservative Þ 5 C; 2.0%; σ 0 G; 4 T; 0 U; Score 19.4; Pred. No. 1 Mismatches 1.2e+03 0 Other, DB 1; Length 21; Indels 0 Gaps

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RESULT 595

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CAAAGTCCTGGGATTACAGGC 21

S

388 CAAAGTGCTGGGATTACAGGC 408

AAH39585; AAH39585 standard; DNA; 21 ₽P

SNP specific upper PCR primer SEQ ID 2381.

14-AUG-2001

(first entry)

AAH39585
AAH39585
ID AAH1
XX
AC AAH1
XX
DT 14-J
DT 14-J
XX
Sing
KW Sing
KW SNPJ
KW SNPJ
KW POLJ
KW POLJ
KW POLJ
XX
XX
PD 26-J
XX
PPD 26-J
XX
XX polycystic kidney disease; osteogenesis acute intermittent porphyria: rhemmatric Single nucleotide Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; inflammation; disease; osteogenesis imperfecta; autoimmune disease; porphyria; rheumatoid arthritis; multiple sclerosis; paternity

Homo sapiens.

26-APR-2001.

13-OCT-2000; 2000WO-US028436.

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RESULT 596
AAH88861/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cc performing a single-nucleotide primer extension reaction. The coligonucleotides are useful for determining the presence, absence or cc identity of a SNP and for genotyping nucleic acid samples, for e.g. to casses by association analysis the genotype of an individual or group of cindividuals, having a pathological phenotypic trait suspected of being cc caused by one or more SNPs. Phenotypic traits include diseases e.g. cc agammaglobulinaemia, diabetes insipidus, Lesch-Nyhan syndrome, muscular cc dystrophy, familial hypercholesterolaemia, polycystic kidney disease, cc disease of which a component is or may be genetic such as autoimmune cc diseases, including, rheumatoid arthritis, multiple sclerosis, cc inflammation, cancer, nervous system diseases and infection by pathogenic confidence in the stigations and cc microorganism. The method is also useful in forensic investigations and cc for a human SNP containing DNA sequence represents a PCR primer specific for a human SNP containing DNA sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2001-290930/30.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Picoult-Newburg L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       includes kits for determining the presence or absence of a SNP, oligonucleotides of the invention. The PCR primers are used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide primer extension (SNPE) primers, and the sequences of regions flanking sites of single nucleotide polymorphisms SNPs. The present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; Page 62; 83pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      acid
                                                                                                                                                                                         Human; single nucleotide polymorphic; SNP; forensic science; paternity testing; phenotypic trait; genetic mapping; animal breeding; plant breeding; ds.
                                                                                                                                                                                                                                                                                                                09-SEP-2004
27-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 21 BP; 5 A; 9 C; 2 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SNP flanking sequence, the SNPE primer is used as a genotyping primer. The oligonucleotides are useful for genotyping a nucleic acid sample by
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                                                                                                                                                                                                                                                                                                                                                                                                                  AAH88861 standard; DNA;
                                                                                                                                     Homo sapiens
Unidentified
                                                                                                                                                                                                                                                                          Human polymorphic oligonucleotide L39064 fragment #1.
                                                                          Key
variation
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                                                                             Location/Qualifiers
                                                        /*tag= a
                                   standard_name= "single nucleotide polymorphism"
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95.2%;
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sed to amplify a
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WO200134840-A2

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RESULT 597
AAH89111
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Best Local &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention relates to human oligonucleotides comprising a single nucleotide polymorphic site (SNP: ANH8979-AAH89219). The present sequence is one such oligonucleotide. The oligonucleotides can be used if forensics, paternity testing, correlation of polymorphisms with phenotypic traits, genetic mapping of phenotypic traits and marker assisted breeding of animals and crop plants
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New polymorphic sites derived from the human genome are useful to determine sites correlating with phenotypic traits, particularly and also in forensics and paternity testing.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10-NOV-2000; 2000WO-US030766.
                                                                                                                                                                                                                            Human; single nucleotide polymorphic; SNP; forensic science; paternity testing; phenotypic trait; genetic mapping; animal plant breeding; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 21 BP; 4 A; 5 C; 6 G; 6 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10-NOV-1999;
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27-FEB-2002
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                                                                                                                                                                                                                                                                              Human polymorphic oligonucleotide U29874 fragment #4.
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                                                                                                                                                                                                                                                                                                                                                                   AAH89111 standard; DNA; 21 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (GLAX ) GLAXO GROUP LTD (AFFY-) AFFYMETRIX INC.
(GLAX ) GLAXO GROUP LTD. (AFFY-) AFFYMETRIX INC.
                                                                                                          WO200134840-A2
                                                                                                                                                         variation
                                                                                                                                                                    ĕеу
                                                                                                                                                                                          Unidentified
                                                                                                                                                                                                      Homo sapiens
                                   10-NOV-1999;
                                                           10-NOV-2000; 2000WO-US030766.
                                                                                   17-MAY-2001.
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ilarity 95.2%;
Conservative
                                                                                                                                                                                                                                                                                                        (revised)
(first entry)
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                                    99US-0164596P.
                                                                                                                                                         Location/Qualifiers
                                                                                                                      /*rag= a
/standard_name= "single nucleotide polymorphism"
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Pred. No. 1.2e+03
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RESULT 59
AAF74150
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Best Local S
Matches 20
serotonin member 4 (SLC6A4) gene or a fragment of it or a sequence recomblementary to the first sequence. The invention is used in producing recombinant organism that can be used to express SLC6A4 for protein structure analysis and binding studies. A commonsition
                                                                                                                                                                                                                           gene for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim
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                                                                                                                                                                                                                                              New isolated polynucleotide comprising a polymorphic variant for the solute carrier family 6 neurotransmitter transporter, serotonin member 4 gene for identifying drugs for treating disorders related to expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Solute carrier family 6 neurotransmiter transporter; genotyping; allele specific oligonucleotide; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAF74150;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New polymorphic sites derived from the humar determine sites correlating with phenotypic and also in forensics and paternity testing.
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                                                                                                                                                                           Example 1; Page
                                                                                                                                                                                                                                                                                                                                                                                               Denton
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                                                                                  present invention relates to a polymorphic variant of a reference uence for the solute carrier family 6 neurotransmitter transporter, otonin member 4 (SLG6A4) gene or a fragment of it or a sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                              GENAISSANCE
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Pred. No. 1.
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nenotypic traits, particular
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.2e+03;
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                                                                                                                                                                                                                                                                                                                                                                                            Stephens
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                                                                                                                                                          allelic variant polynucleotide. The polypeptides of the invention have neuroleptic activity. The polynucleotides may have a use in gene therapy. DISC1 or DISC2 nucleic acid molecules are useful for disgnosting or treating a subject having a disease or disorder associated with specific DISC1 or DISC2 alleles and/or abstrant DISC1 expression or activity e.g. neuropsychiatric disorder such as schizoaffective, bipolar, unipolar affective or adolescent conduct disorder or schizophrenia. Similarly, the
                                                                                                                         compound that inhibits DISC1 protein activity may be used in the method for treating such neuropsychiatric disorders. The sequences shown in ABQ93575-ABQ93658 represent the PCR primers used in the invention to
                                                                                                                                                                                                                                                                                      New human Disrupted-In-Schizophrenia (DISC) 1 and DISC2 single nucleotide polymorphisms, useful for preventing neuropsychiatric disorders e.g. schizophrenia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       genotyping oligonucleotide
                                                                                         Sequence
                                                                                                                                                                                                                                                                Claim 17; Fig 4; 169pp;
                                                                                                                                                                                                                                                                                                                                      WPI; 2002-590791/63.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human, Disrupted In Schizophrenia 1; DISC1; neuroleptic; gene therapy; neuropsychiatric disorder; schizoaffective disorder; bipolar disorder; unipolar affective disorder; adolescent conduct disorder; schizophrenia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABQ93614;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
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                                                                                                                                                                                                                                                                                                                                                            Meyer JM,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human DISC1/DISC2 PCR primer
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                                                       Similarity
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                                            Conservative
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RESULT 600
Caryl hydrocarbon (AHR), aryl hydrocarbon receptor nuclear translocator (C (ARNT), cathepsin S (CTSS), cyclooxgenase 2 (COX2), diazepam binding C (inhibitor (DBI), epoxide hydroxylase 2 (EPHX2), 5-lipoxygenase activating C protein (FLAP), glutathione-S-transferase 12 (GST12), histamine-N-methyl C transferase (HNWT), (Kallikrein 2) KLK2, nicotinamide -N-methyl C transferase (NNWT), MADPH quinnone oxidoreductase 2 (NQC2), c transferase (NNWT), NADPH quinnone oxidoreductase 2 (NQC2), c transferase (NUNT), NADPH quinnone oxidoreductase 2 (NQC2), c transferase (NCTSB1), UDP-glucuronosyl transferase 2B4 (UGT2B4), UDP-glucuronosyl transferase 2B7 (UGT2B7), UDP-glucuronosyl c transferase (UGT2B1), urokinase receptor (NPA), multidrug resistance 1 (MRP3), lactotransferrin (LTF), multidrug resistance associated protein 3 (MRP3), orphan nuclear receptor (NR112), or acetylcholine muscarinic C receptor 1, 2, 3, 4, or 5 (CHMR1, CHMR2, CHMR3, CHMR4 or CHMR5) sequence. C The polymorphisms in the human genes cited in the invention are useful as C genetic linkage markers for locating and characterising the genes that C are responsible for specific traits within the genome and eventually cidentifying the genes responsible for a variety of disorder-related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Whuman; ds; cytochrome P450 A1; CYP4501A1; UGT2B4; MDR1;
W cytochrome P450 A2; CYP4501A2; cytochrome P450 02E; CYP45002E1; LTF;
W adrenergic receptor beta1; ADBR1; arryl hydrocarbon; ARN; MRP3; NR112;
W aryl hydrocarbon receptor nuclear translocator; ARNT; cathepsin S; CTSS;
W cyclooxgenase 2; COX2; diazepam binding inhibitor; DBI; haematological;
W cyclooxgenase 2; EPHX2; 5-lipoxygenase activating protein; FLAP;
W cyclooxgenase 2; EPHX2; 5-lipoxygenase activating protein; FLAP;
W cyclooxgenase 2; KLK2; nicotinamide-N-methyl transferase;
W HNMT; kallikrein 2; KLK2; nicotinamide-N-methyl transferase; NNMT;
W NADPH quinone oxidoreductase 2; NQO2; sulfotransferase thermolabile; STM;
W NADPH quinone oxidoreductase 21, NQO2; sulfotransferase thermolabile; STM;
W UDP-glucuronosyl transferase 2B4; UDP-glucuronosyl transferase 2B7;
W UGT2B7; UDP-glucuronosyl transferase; UGT2B15; urokinase receptor; uPA;
W multidrug resistance 1; lactotransferrin; orphan nuclear receptor;
W multidrug resistance associated protein 3; cancer; prostate;
W multidrug metabolism; cardiovascular function; colorectal tumour;
W actelylcholine muscarinic receptor; publiconary; immunological; SNP;
W central nervous system; pulmonary; immunological; SNP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human multidrug resistance gene polymorphic sequence #66
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                                                                                                                                                                                                                                                                                                                                                                                                                                 This invention relates to the sequence of an isolated nucleic acid molecule comprising at least one base variation from that of a known human cytochrome P450 A1 (CYP4501A1), cytochrome P450 A2 (CYP4501A1), adrenergic receptor betal (ADBR1), cytochrome P450 02E1 (CYP4500ZE1), adrenergic receptor betal (ADBR1),
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Isolated nucleic acid molecules having polymorphisms in known human genes e.g. cytochrome p450 and cathepsin S useful as genetic linkage markers for locating, identifying and characterizing the genes responsible for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200257410-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           single nucleotide polymorphism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        22; Page 144; 714pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hall J;
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RESULT 601
ADC42593
ID ADC425
XX ADC425
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XX Cances
CX Chemos
XX Cances
KW Chemos
XX Synth
XX CACS
PN WO200:
XX WD1;
XX Diagn
PH Diagn
PH Diagn
PH Cance
XX WPI;
XX Diagn
PH Cance
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                                                                                                                                                                                                                                                                                    Diagnosing or determining cancer or increased risk of cancer in a patient, by testing Fanconi Anemia/BRCA pathway gene or protein for a cancer-associated defect, that indicates cancer or increased risk of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 21 BP; 5 A; 5 C; 7 G; 4 T; 0 U; 0 Other;
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02-NOV-2001; 2001WO-US045561.
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The invention relates to a novel method of diagnosing or determining if patient has cancer or is at increased risk of cancer, involving testing Fanconi Anaemia (FA)/BRCA pathway gene or protein for the presence of a cancer-associated defect, where the presence of one or more cancer-associated defects is indicative of cancer or an increased risk of cancer associated defects is indicative of cancer or an increased risk of cancer associated defects is indicative of cancer or an increased risk of cancer associated defects is indicative of cancer or an increased risk of cancer associated defects is indicative of cancer or an increased risk of cancer associated defects is indicative of cancer or an increased risk of cancer.

an increased risk of cancer as cytostatic activity. The

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Example 14;

Page 101; 160pp; English

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The invention relates to an isolated nucleic acid (N1) comprising at least 20 but not more than 1500 consecutive nucleotides of the optineurin promoter appearing as ADE13890. Also included are the optineurin promoter operably linked to a heterologous nucleic acid, a nucleic acid capable of detecting a single nucleotide polymorphism (SNP) in the optineurin promoter, a host cell comprising the promoter operably linked to a heterologous sequence, diagnosing or promoter a sequence, diagnosing or promoter a cell or bodily fluid (comprising detecting a polymorphism in a promoter region of the optineurin gene, associated with a glaucoma phenotype), detecting a SNP sequence variation in a sample containing
                                                                                                                                                                                                                                                                                                                                                                                                                                                 New nucleic acid sequences of the optineurin gene are useful to detect polymorphisms particularly single nucleotide polymorphisms in the optineurin promoter to diagnose, prognose and treat glaucoma and related
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The present sequence represents a non-nucleotide probe. The probe is useful for suppressing the binding of one or more detectable nucleic acid probes, that are greater than 100 base pairs and that have been derived from genomic nucleic acid, to one or more undesired sequences in an assay for determining target genomic nucleic acid of a sample. The method comprises contacting the sample with the mixture of probes (preferably comprising 5-50 probes), contacting the sample with the one or more detectable nucleic acid probes, and determining the target genomic nucleic acid of the sample by determining the hybridization of the one or more detectable nucleic acid probes to the target genomic nucleic acid of the sample by determining the hybridization of the one or

Claim 10; SEQ ID NO 25; 103pp; English

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RESULT 603
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                                                                                                                                                                                                                                                                                                                                                                                                                 Non-nucleotide probe for suppressing binding of detectable nucleic acid probes to undesired sequences, has aggregate nucleobase sequence homologous to randomly distributed repeat sequence of genomic nucleic
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CC the sample. The genomic nucleic acid is contained in a fixed tissue or a CC cell, and the sample is metaphase spreads, interphase nucleic or nucleic found in paraffin embedded tissue material or frozen tissue sections. The CC probe is also useful in comparing a sample of genomic nucleic acid with the method comprises treating a sample of genomic nucleic acid with CC control genomic nucleic acid, which are differentially labelled, the CC array or both the sample and control genomic nucleic acid and the array with treated mixture of sample and control genomic conditions, CC contacting the array with treated mixture of sample and control genomic cut to the signals from the differential labels of the array to the signals from the differential labels of the array to the caused by hybridization of the probes to genomic nucleic acid, thus determining one or more variations in copy numbers of sequences in the control genomic nucleic acid with the relative copy numbers of substantially complete as compared with the relative copy numbers of substantially complete as compared with the relative copy numbers of substantially complete as compared with the relative copy numbers of substantially complete as compared with the relative copy numbers of substantially complete as assumed to genomic nucleic acid to be tested and the reference of nucleic acid are labelled with detectable molety such that hybridization of the genomic array is determined by determining the presence, absence, amount or location of the detectable label on the one or more genomic constant and the reference of anount or location of the detectable and the reference, absence, amount or location of the detectable label on the one or more genomic constants and comparises nucleic acid that is prepared from the one or more genomic constants and comparises nucleic acid that is prepared from the one or more genomic constants and comparises nucleic acid that is prepared from the one or more genomic constants and comparises nucleic acid that is prepared from the 
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ADH59612/c
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Best Local Similarity
Matches 20; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   non-nucleotide probe; Bacterial Artificial Chromosome clone; BAC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Non-nucleotide probe of the invention #16.
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Non-nucleotide probe for suppressing binding of detectable nucleic acid probes to undesired sequences, has aggregate nucleobase sequence homologous to randomly distributed repeat sequence of genomic nucleic
                                                                                                                                                              WPI; 2003-421160/39.
                                                                                                                                                                                                                                               Kirtsen NV,
                                                                                                                                                                                                                                                                                                                                                                  (BOST-)
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DAKOCYTOMATION DENMARK AS
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Pred. No. 1.2e+03;
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Claim 10; SEQ ID NO 18; 103pp; English

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  RESULT 605
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 21 BP; 4 A; 6 C; 8
                                                                                                                                                                                                                                                                                                                     non-nucleotide probe; Bacterial Artificial Chromosome clone; BAC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       represents a non-nucleotide
                                                                    24-SEP-2001;
                                                                                                                  24-SEP-2002; 2002WO-US030573
                                                                                                                                                                                                               WO2003027328-A2.
                                                                                                                                                                                                                                                        Synthetic
                                                                                                                                                                                                                                                                                                       probe.
                                                                                                                                                                                                                                                                                                                                                                         Non-nucleotide probe
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(BOST-) EOSTON PROBES INC. (DAKO-) DAKOCYTOMATION DENMARK AS
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                                                                      2001US-0324499P
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CC probes, that are greater than 100 base pairs and that have been derived
CC from genomic nucleic acid, to one or more undesired sequences in an assay
CC for determining target genomic nucleic acid of a sample. The method
CC comprising 5-50 probes), contacting the mixture of probes (preferably
CC comprising 5-50 probes), contacting the mixture of probes (preferably
CC comprising 5-50 probes), contacting the sample with the one or more
CC detectable nucleic acid probes, and determining the target genomic
CC uncleic acid of the sample by determining the hybridization of the one or
CC more detectable nucleic acid probes to the target genomic nucleic acid of
CC the sample. The genomic nucleic acid is contained in a fixed tissue or a
CC cell, and the sample is metaphase spreads, interphase nucleic acid with
CC that of a control sample using a sample of genomic nucleic acid with
CC that of a control sample using a sample of genomic nucleic acid with
CC control genomic nucleic acid, which are differentially labelled, the
CC contacting the array with treated mixture of sample and control genomic
CC contacting the array with treated mixture of sample and control genomic
CC contacting one or more variations in copy numbers of sequences in the
CC identical sequences in the control. The hybridization conditions,
CC determining one or more variations in copy numbers of substantially
CC array is determined using an intercalating dye or a detectable antibody,
CC or its fragment, that is specific for a nucleic acid hybrid.
CC array is determined using an intercalating the presence, absence,
CC arrays. The genomic array is determined by determining the presence, absence,
CC arrays to the detectable nucleic acid hybrid.
CC arrays to the detectable nucleic acid the reference of more genomic
CC arrays to the detectable nucleic acid the reference, absence,
CC arrays to the detectable label on the one or more genomic
CC arrays to the detectable nucleic aci
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ADH59617/c
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Matches 20
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Synthetic
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                                                                                                                                                                                                                                                 ADH59617 standard; DNA; 21 BP
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                                                                 non-nucleotide probe; Bacterial Artificial Chromosome clone; BAC;
                                                                                                           Non-nucleotide probe of the invention #21.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            arrays. The genomic array comprises nucleic acid that is prepared from Bacterial Artificial Chromosome (BAC) clones. The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
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                                                                                                                                                          (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 19.4; DB 1;
Pred. No. 1.2e+03;
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ADH59605

ADH59605 standard; DNA; 21 BP

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cc array or both the sample and control genomic nucleic acid and the array control genomic nucleic acid and the array with the sample and control genomic nucleic acid and the array with treated mixture of sample and control genomic nucleic acid under suitable hybridization conditions, and comparing the content acid of the signals from the differential labels of the array to that caused by hybridization of the probes to genomic nucleic acid, thus containing one or more variations in copy numbers of sequences in the control. The hybridization of the genomic content array to determined using an intercalating dye or a detectable antibody, cor its fragment, that is specific for a nucleic acid/nucleic acid hybrid. The sample of genomic nucleic acid to be tested and the reference of nucleic acid are labelled with detectable moiety such that hybridization of the genomic array is determined by determining the presence, absence, armount or location of the detectable holety such that hybridization of the genomic array is determined by determining the presence, absence, armount or location of the detectable noiety such that hybridization of the genomic array comprises nucleic acid that is prepared from CC arrays. The genomic array comprises nucleic acid that is prepared from CC arrays. The genomic acray comprises nucleic acid that is prepared from CC arrays. The genomic acray comprises nucleic acid that is prepared from CC arrays. The genomic acray comprises nucleic acid that is prepared from CC arrays. The genomic acray comprises nucleic acid that is prepared from CC arrays.
Query Match
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Matches 20
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                                                                                                           Sequence
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DAKOCYTOMATION DENMARK AS
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  2.0%;
nilarity 95.2%;
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                                                                                                              A; 5 C; 8
     0;
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                               Score 19.4;
Pred. No. 1.
        Mismatches
                                  1.2e+03
                                                        DB 1; Length 21;
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comprises contacting the sample with the mixture of probes (preferably comprising 5-50 probes), contacting the sample with the one or more detectable nucleic acid probes, and determining the target genomic common detectable mucleic acid probes, and determining the target genomic common detectable mucleic acid probes to the target genomic nucleic acid of the sample. The genomic nucleic acid is contained in a fixed tissue or a control genomic nucleic acid is contained in a fixed tissue or a control genomic nucleic acid is contained in a fixed tissue or a control genomic nucleic acid in the sample of genomic nucleic acid with that of a control sample using a genomic nucleic acid reference array. The method comprises treating a sample of genomic nucleic acid and control genomic nucleic acid, which are differentially labelled, the array or both the sample and control genomic nucleic acid and control genomic nucleic acid under suitable hybridization conditions, contacting the array with treated mixture of sample and control genomic nucleic acid with the mixture of the signals from the differential by and comparing the control sample as compared with the relative copy numbers of sequences in the control. The hybridization of the genomic determined using an intercalating dye or a detectable antibody, or its fragment, that is specific for a nucleic acid/nucleic acid hybrid. The sample of genomic nucleic acid to be tested and the reference of nucleic acid and the reference of controls array is determined by determining the presence, absence, absence, absence, and acceptable antibody.

CC arrays. The genomic array comprises mucleic acid that is prepared from conditions of the detectable mother one or more genomic conditions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             non-nucleotide probe; Bacterial Artificial Chromosome clone;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Non-nucleotide probe of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADH59605;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 probes, that are greater than 100 base pairs and that have been determine mucleic acid, to one or more undesired sequences in an for determining target genomic nucleic acid of a sample. The method comprises contacting the sample with the mixture of probes (preferal nore)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Non-nucleotide probe for suppressing binding of detectable nucleic acid probes to undesired sequences, has aggregate nucleobase sequence homologous to randomly distributed repeat sequence of genomic nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                useful for suppressing the binding of one or more detectable nucleic ac probes, that are greater than 100 base pairs and that have been derived from genomic nucleic acid, to one or more undesired sequences in an ass
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2003-421160/39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  24-SEP-2002; 2002WO-US030573
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present sequence represents a non-nucleotide probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 10; SEQ ID NO 11; 103pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       24-SEP-2001; 2001US-0324499P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AS
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Query Match

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non-nucleotide A;

3AC) clones. The present of the invention.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             XW x-linked anhidrotic ectodermal dysplasia; al incontinentia pigmenti; XW influenza; rheumatoid arthritis; influenzarony bowel disease; colltis; XW atherosclerosis; cachexia; euthyroid sick syndrome; stroke; EAE; XW experimental allergic encephalomyelitis; autoimmune disorder; wound; XW hyper immune activity; acute phase response; hypercongenital condition; XW birth defect; necrotic lesion; organ transplant rejection; pancreas; XW signal transduction; hyperproliferative disorder; diabetes mellitus; XW vitamin B12 malabsorption; neurological disorder; Huntington's chorea; XW vitamin B12 malabsorption; neurological disorder; Huntington's chorea; XW unner's syndrome; bacterial infection; cardiovascular disorder; XW unfertility; psoriasis; haemolytic anaemia; antiinflammatory; anti-HIV; XW cytostatic; hepatotropic; virucide; antinheumatic; antiarthritic; XW untiasthmatic; immunosuppressive; vulnerary; antibacterial; XW antiasthmatic; immunosuppressive; vulnerary; antibacterial; XW antiafertility; antianaemic; antispsoriatic; cerebroprotective; cardiant; XW antiarteriosclerotic; PCR; primer; ss.
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The present invention relates to the isolation of human nuclear factor. RappaB (NF-kappaB) associated polypeptides and polynucleotides. The NF-kappaB associated polypeptide and polynucleotide sequences are useful if preventing, treating or ameliorating various disorders including immune disorders, inflammatory disorders, cancers, disorders relating to aberrant apoptosis, hepatic disorders, hodgkin's lymphomas, haematopoietic tumours, hyper-igm syndromes, hypohidrotic ectodermal dysplasia, rimmunodeficiency, viral infections (e.g. those caused by human immunodeficiency virus (HIV), human T-cell lymphotropic virus (HTLV), hepatitis B, hepatitis C, Epstein Barr virus (EBV), influenza),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 inflammatory disorder; apoptosis; hepatic disorder; Hodgkin's lymphom
haematopoietic tumour; hyper-IgM syndrome; viral infection; asthma;
hypohidrotic ectodermal dysplasia; human immunodeficiency virus; HIV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19-APR-2001; 2001US-0284962P
26-APR-2001; 2001US-0286645P
09-JAN-2002; 2002US-0346986P
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                                                                                                                                                                                                                                                                                                               diagnosing,
                                                                                                                                                                                                                                                                                                                 Novel NF-kappaB-associated polypeptides and polynucleotides useful for diagnosing, treating and preventing cancer, hepatic disorders, aberrant
                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-093119/08.
                                                                                                                                                                                                                                                                                                                                                                                                                    Carman J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                            (BRIM ) BRISTOL-MYERS SQUIBB CO
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                                                                                                                                                                                                                                                     Example 3;
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                                                                                                                                                                                                                                                                                                                                                                                                                    Feder J,
                                                                                                                                                                                                                                                     Page 341; 608pp; English.
                                                                                                                                                                                                                                                                                                viral infections, autoimmune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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                                                                                                                                                                                                                                                                                                  disorders,
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                                                                                                                                                                                                              nuclear factor-
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                                                                                                                                                                                                                                                                                                  Novel isolated LPDL or LPDLR lipase polypeptides, useful for substances that bind to the protein and which are useful for diseases associated with lipase function e.g. atherosclerosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21-DEC-2001; 2001US-0341786P
10-JAN-2002; 2002US-0346603P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         lipase; LPDL; LPDLR; lipase deficiency; atherosclerosis; fatty liver disease; dyslipidaemia; hypercholesterolaemia; hypertriglyceridaemia; mixed dyslipidaemia, lipid deficient state; hypertriglyceridaemia; state; human; ss; PCR; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human LPDLR PCR primer #12.
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TSUI L.
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                                                                                                                                                                                                                                                                                                                                  for identifying for treating
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The invention relates to an isolated mammalian (e.g., human or mouse) lipase polypeptide (polyp), e.g., LPDL (I) or LPDLR polyp (II). (I) or (II) 'is useful for identifying substances which can bind with LPDL or LPDLR polyp, and for identifying a compound that affects the binding of LPDL or LPDLR polyp and an LPDL or LPDLR binding polyp. (I) or (II) or their nucleic acid is useful for identifying a compound that affects LPDL or LPDLR polyP activity or expression. (I) or (II) or their nucleic acid

Disclosure;

SEQ ID NO 68; 172pp; English

hypercholesterolemia.

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RESULT 610
ADO12329
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Best Local
The invention relates to a method of designing primers for simultaneous amplification of target DNA fragments in a single multiplex polymerase chain reaction by aligning a first primer and a second primer; The method comprises: (a) aligning a first primer and a second primer; and (b) selecting the first primer where the first primer at its 3' end does not contain four or more bases that are perfectly matching to the 3' end sequence of the first primer or a second primer, the first primer at its 3' end does not contain seven or more bases that are perfectly matching except one mismatch to the 3' end sequence of the first primer or the second primer, the first primer at its 3' end does not contain save or the first primer at its 3' end does not contain sav or the second primer, the first primer at its 3' end does not contain sax or the second primer, the first primer at its 3' end does not contain sax or the second primer, the first primer at its 3' end does not contain sax or the second primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 increased or decreased LPDL or LPDLR expression or activity in an animal, where the condition is lipse deficiency, atherosclerosis, fatty liver disease and dyslipidemias, such as hypercholesterolemia, hypertriglyceridemia, mixed (combined) dyslipidemia, lipid or lipoprotein deficient states, and/or any other tissue or plasma disorders of lipid or lipoprotein metabolism. The nucleic acid is useful for diagnosing the presence of or a predisposition for a disorder in a subject which involves detecting a germline alteration in the nucleic acid in the subject an inhibitor is useful for modulating trajlyceride activity by inhibiting expression or activity of (I) or (II). The nucleic acid is useful as a probe or primer. The present sequence is used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ss; primer; simultaneous amplification; single multiplex polymerase chain reaction; multifactorial genetic alteration; pharmacogenetic reaction; genotyping:
                                                                                                                                                                                                                                                                              Designing primers for simultaneous amplification of target DNA fragments in a single multiplex polymerase chain reaction, for high throughput
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         07-OCT-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Single multiplex PCR primer #1701
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADO12329;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          exemplification of the invention.
                                                                                                                                                                                                                                                                 multiplex DNA
                                                                                                                                                                                                                                                                                                                                          WPI; 2004-340914/31.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     07-OCT-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO2004033649-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADO12329
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 is useful for detecting or monitoring a condition associated with
                                                                                                                                                                                                                                                                                                                                                                                                                  (UYNE-) UNIV NEW JERSEY MEDICINE & DENTISTRY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       expression
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TGCTGGGATTACAGGCGTGAG 883
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TGCTGGGATTACAGGCATGAG 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 5
                                                                                                                                                                                                                            Page 41; 120pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                                                                 sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA;
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                                                                                                                                                                                                                                                                 amplification,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 19.4;
Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
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                                                                                                                                                                                                                                                                 comprises aligning
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              .2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       genotyping; polymorphism;
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                                                                                                                                                                                                                                                                   two primers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              disease;
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                                                                                                                                                       The method
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RESULT 611
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    first primer or the second primer, and the first primer at its 3' end does not contain eleven or more bases that are perfectly matching except one mismatch to a sequence anywhere of the first primer or the second primer. The method is useful for designing primers for simultaneous amplification of target DNA fragments in a single multiplex polymerase chain reaction. It is also useful in the identification of multiple genes related to multifactorial diseases, the genome-scale detection of genetic alterations, the studies in pharmacogenetic reactions, the genotyping genetic polymorphisms in a large population, the gene expression profiling in various samples and high throughput genotyping technologies. This sequence corresponds to an example of a primer of the invention.
                                                                The present invention describes a restriction primer for eukaryotic short interspersed repeated sequences (SINE), which has one or more additional bases that are a mismatch to, or are unrelated to, the 3'-terminal end of the SINE. The annealing temperature of the primer to the DNA sequence is kept higher than the fusion temperature of the primer during polymerase chain reaction (PCR). The PCR fragments obtained are subjected to electrophoresis to obtain a fingerprint. By comparing the polymorphs from the electrophoresis band pattern, eukaryotic individuals are distinguished. The primer is used for amplifying a eukaryotic deoxyribonucleic acid (DNA) sequence, pinched between SINE sequences by polymerase chain reaction (PCR) fingerprinting. In particular it may be used individual identification of humans for medical and legal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; short interspersed repetitive element; SINE; PCR; primer; Oncorhynchus; restriction primer; short interspersed repeated sequence; eukaryote; restricted polymerase chain reaction fingerprinting; identification; DNA specimen; discrimination; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human short interspersed repetitive element PCR primer #14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAZ25156 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Restriction primer for distinguishing individuals with short interspersed repeated sequence of eukaryotes by restricted polymerase chain reaction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13-DEC-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 6; Page 3; 17pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (NORQ ) NORINSUISANSHO SUISANCHO YOSHOKU KENKYUSHOCHO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
used individual identification of humans for medical applications and ecological studies. DNA specimens in (approximately 10 ng in mass) can be used for individuals.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP; 4 A; 8 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      98JP-00195692.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2.0%;
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Pred. No. 1.
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for individual

discrimination

Best Local Query Match

Similarity

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Score 19.4; Pred. No. 1.

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ID AAZ251
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                                     The present invention describes a restriction primer for eukaryotic short characteristic interspersed repeated sequences (SINE), which has one or more additional bases that are a mismatch to, or are unrelated to, the 3'-terminal end of the SINE. The annealing temperature of the primer to the DNA sequence is chain reaction (PCR). The PCR fragments obtained are subjected to celectrophoresis to obtain a fingerprint. By comparing the polymorphs from the electrophoresis band pattern, eukaryotic individuals are decoxyribonucleic acid (DNA) sequence, pinched between SINE sequences by colymerase chain reaction (PCR) fingerprinting, In particular it may be used individual identification of humans for medical and legal capplications and ecological studies. DNA specimens in traces applications and ecological studies. DNA specimens in traces of eukaryotes using the primer in a polymerase chain reaction (PCR). Co feukaryotes using the primer in a polymerase chain reaction (PCR).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       of eukaryotes using the primer in a polymerase chain reaction (PCR) AAZ25143 to AAZ25191 represent specifically claimed examples of printrom the present invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Oncorhynchus; restriction primer; short interspersed repeated sequence; enharyote; restricted polymerase chain reaction fingerprinting;
                                                                                                                                                                                                                                                                                                                                                        Restriction primer for distinguishing individuals with short interspersed repeated sequence of eukaryotes by restricted polymerase chain reaction
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human short interspersed repetitive element PCR primer #26.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAZ25168 standard;
                                                                                                                                                                                                                                                                                                            Claim 6; Page 4; 17pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10-JUL-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; short interspersed repetitive element; SINE; PCR; primer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              identification;
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                           present
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA specimen; discrimination; ss.
                              invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                           SUISANCHO YOSHOKU KENKYUSHOCHO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 19.4;
Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
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U;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length
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                                                                                                                               RESULT 613
AAZ25162
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밁
                                                             Query Match
Best Local S
Matches 20
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                                                                                                                                                                      the electrophoresis band pattern, eukaryotic individuals are distinguished. The primer is used for amplifying a eukaryotic destyribonucleic acid (DNA) sequence, pinched between SINE sequences by polymerase chain reaction (PCR) fingerprinting. In particular it may be used individual identification of humans for medical and legal applications and ecological studies. DNA specimens in traces (approximately 10 ng in mass) can be used for individual discrimination of eukaryotes using the primer in a polymerase chain reaction (PCR).

AAZ25143 to AAZ25191 represent specifically claimed examples of primers
                                                                                                                                                                                                                                                                                                      The present invention describes a restriction primer for eukaryotic short interspersed repeated sequences (SINE), which has one or more additional bases that are a mismatch to, or are unrelated to, the 3'-terminal end of the SINE. The annealing temperature of the primer to the DNA sequence is kept higher than the fusion temperature of the primer during polymerase chain reaction (PCR). The PCR fragments obtained are subjected to electrophoresis to obtain a fingerprint. By comparing the polymorphs from the electrophoresis band pattern, eukaryotic individuals are
                                                                                                                             Sequence 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Restriction primer for distinguishing individuals with short interspersed repeated sequence of eukaryotes by restricted polymerase chain reaction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28-JUN-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Oncorhynchus; restriction primer; short interspersed repeated sequence; eukaryote; restricted polymerase chain reaction fingerprinting; identification; DNA specimen; discrimination; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; short interspersed repetitive element; Oncorhynchus; restriction primer; short inters
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human short interspersed repetitive element PCR primer #20.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAZ25162 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1999-583348/50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10-JUL-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        13-DEC-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          fingerprinting.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (NORQ ) NORINSUISANSHO SUISANCHO YOSHOKU KENKYUSHOCHO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       sapiens
                              869
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20;
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                                                              20;
                                                                             Similarity
                  GATTACAGGCGTGAGCCACCA 889
                                                                                                                                                             present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Page 3; 17pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GATTACAGGCGTGAGCCACTA 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GATTACAGGCGTGAGCCACCA 889
                                                                                                                           BP; 6 A; 5 C; 8 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first
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                                                                            2.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      22
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                                                             0
                                                                            Score 19.4;
Pred. No. 1.
                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                            .3e+03
                                                                                             DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SINE; PCR; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1;
                                                                                           Length 22;
                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
                                                             0;
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                                                             Gaps
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RESULT 614

BXH

13-DEC-1999

(first entry)

Human short interspersed repetitive element PCR primer #25

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RESULT 615
AAZ25167
ID AAZ251
XX
AC AAZ251
XX
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ID AAZ2
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                                                                                                                                                                                                                                                             The present invention describes a restriction primer for eukaryotic short CC interspersed repeated sequences (SINE), which has one or more additional CC bases that are a mismatch to, or are unrelated to, the 3'-terminal end of CC the SINE. The annealing temperature of the primer to the DNA sequence is kept higher than the fusion temperature of the primer during polymerase CC chain reaction (PCR). The PCR fragments obtained are subjected to electrophoresis to obtain a fingerprint. By comparing the polymorphs from CC distinguished. The primer is used for amplifying a eukaryotic deoxyribonucleic acid (DNA) sequence, pinched between SINE sequences by CC polymerase chain reaction (PCR) fingerprinting. In particular it may be used individual identification of humans for medical and legal CC applications and ecological studies. DNA specimens in traces CC elekaryotes using the primer in a polymerase chain reaction (PCR).

CC ALZZ5143 to ALZ25191 represent specifically claimed examples of primers
                                                                                                                                                                                                 Matches
                                                                                                                                                                                                            Query Match
Best Local Similarity
                  AAZ25167;
                                                AAZ25167 standard; DNA;
                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 6; Page 3; 17pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Restriction primer for distinguishing individuals with short interspersed repeated sequence of eukaryotes by restricted polymerase chain reaction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Oncorhynchus; restriction primer; short interspersed repeated sewharyote; restricted polymerase chain reaction fingerprinting;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human short interspersed repetitive element PCR primer #19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAZ25161;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAZ25161 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    tingerprinting.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1999-583348/50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               10-JUL-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             identification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; short interspersed repetitive element; SINE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (NORQ ) NORINSUISANSHO SUISANCHO YOSHOKU KENKYUSHOCHO
                                                                                                                                                              869
                                                                                                                                                                                               20;
                                                                                                                                _
                                                                                                                                GATTACAGGCGTGAGCCACGA 21
                                                                                                                                                              GATTACAGGCGTGAGCCACCA 889
                                                                                                                                                                                                                                                             22 BP; 7
                                                                                                                                                                                                                                                                                           present invention
                                                                                                                                                                                               Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               98JP-00195692
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA specimen; discrimination; ss.
                                                                                                                                                                                                                                                             A; 5 C; 7
                                                                                                                                                                                                            2.0%;
                                                   ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ВP
                                                                                                                                                                                             0,
                                                                                                                                                                                                                                                             G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                            Score 19.4;
Pred. No. 1.
                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            interspersed repeated sequence;
                                                                                                                                                                                                         .3e+03;
                                                                                                                                                                                                                             DB 1;
                                                                                                                                                                                                                           Length 22
                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PCR; primer;
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                                                                                                                                                                                               Gaps
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AAZ25155
ID AAZZ
XX
AC AAZZ
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AC AAZZ
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DT 13-1
XX
DHuma
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DH Huma
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Hum
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Onco
KW Onco
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KW Ouk
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OS Syn
OS Syn
OS Syn
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention describes a restriction primer for eukaryotic short clinterspersed repeated sequences (SINE), which has one or more additional CC bases that are a mismatch to, or are unrelated to, the 3'-terminal end of the SINE. The annealing temperature of the primer to the DNA sequence is CC kept higher than the fusion temperature of the primer during polymerase CC chain reaction (PCR). The PCR fragments obtained are subjected to CC electrophoresis to obtain a fingerprint. By comparing the polymorphs from CC distinguished. The primer is used for amplifying a eukaryotic contained are subjected to CC deoxyribonucleic acid (DNA) sequence, pinched between SINE sequences by CC polymerase chain reaction (PCR) fingerprinting. In particular it may be used individual identification of humans for medical and legal CC applications and ecological studies. DNA specimens in traces CC (approximately 10 ng in mass) can be used for individual discrimination CC eukaryotes using the primer in a polymerase chain reaction (PCR). CC ARZ25143 to ARZ25191 represent specifically claimed examples of primers
                                                                                                                                                                                                                                                                                                                 RESULT
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
Matches 20; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 6; Page 4; 17pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Restriction primer for distinguishing individuals with short interspersed repeated sequence of eukaryotes by restricted polymerase chain reaction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10-JUL-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; short interspersed repetitive element; SINE; PCR; primer; Oncorhynchus; restriction primer; short interspersed repeated sequence; enkaryote, restricted polymerase chain reaction fingerprinting; enkaryote, restricted polymerase chain reaction fingerprinting; identification; DNA specimen; discrimination; ss.
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                                                      eukaryote; restriction primer; short interspersed repeated sequence; eukaryote; restricted polymerase chain reaction fingerprinting; identification; DNA specimen; discrimination.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 22 BP; 7 A; 5 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             fingerprinting.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (NORQ ) NORINSUISANSHO SUISANCHO YOSHOKU KENKYUSHOCHO
                      Synthetic
                                                                                                 Human; short interspersed repetitive element; SINE; PCR; primer; Oncorhynchus; restriction primer; short interspersed repeated so
                                                                                                                                                                                                13-DEC-1999
                                                                                                                                                                                                                                                                          AAZ25155 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                 616
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                                                                                                                                                                                                                                                                                                                                                                                                              698
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                                                                                                                                                                                                                                                                                                                                                                         GATTACAGGCGTGAGCCACTA 21
                                                                                                                                                                                                                                                                                                                                                                                                              GATTACAGGCGTGAGCCACCA 889
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                           interspersed repetitive element PCR primer #13
                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         98JP-00195692
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      95.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2.0%;
                                                                                                                                                                                                                                                                              ΒP
                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 19.4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
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RESULT 617
AAS11629
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CC deproximately 10 ng in mass) can be used for individual discrimination (PCR) in mass chain reaction (PCR)
                                                                                                                                                                          Query Match
Best Local
                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Restriction primer for distinguishing individuals with short interspersed repeated sequence of eukaryotes by restricted polymerase chain reaction
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 6; Page 3; 17pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10-JUL-1998;
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                                                                                                                                                                                                                               Sequence 22 BP; 8 A; 5 C; 6 G;
AAS11629 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            fingerprinting.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (NORQ ) NORINSUISANSHO SUISANCHO YOSHOKU KENKYUSHOCHO
                                                                                                                                                                        Local
                                                                                                                                                                                                                                                                                   eukaryotes using the primer in a polymerase chain reaction (PCR).
Z25143 to AAZ25191 represent specifically claimed examples of primers
                                                                                                                869 GATTACAGGCGTGAGCCACCA 889
                                                                                      Н
                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                  present invention
                                                                                      GATTACAGGCGTGAGCCACAA 21
                                                                                                                                                             Conservative
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DNA;
                                                                                                                                                                            2.0%;
22
 ΒP
                                                                                                                                                             0;
                                                                                                                                                                             Score 19.4;
Pred. No. 1.
                                                                                                                                                                                                                                 3 T;
                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                   o
U;
                                                                                                                                                                               .3e+03
                                                                                                                                                                                                                                   0 Other;
                                                                                                                                                                                                DB 1;
                                                                                                                                                                                              Length
                                                                                                                                                                Indels
                                                                                                                                                                                                  22;
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                                                                                                                                                               Gaps
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CYP2B6; cytostatic; gene therapy; genotyping; cancer; metabolism; human; cancer susceptibility; environmental carcinogen;
                  09-FEB-2000; 2000EP-00102701
                                                                                                           sequencing
                                                                                                                                                Human CYP2B6 allele
                                                                                                                                                                  24-OCT-2001
                                                                                                                                                                                     AAS11629;
                                    09-FEB-2001; 2001WO-EP001456
                                                       16-AUG-2001.
                                                                        WO200159152-A2
                                                                                          Homo sapiens
(EPID-) EPIDAUROS BIOTECHNOLOGIE
                                                                                                            primer
                                                                                                                                                                  (first entry)
                                                                                                                                               sequencing primer seqCYP2B6-7F for exon
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Homo sapiens

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AAF93028/c
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Best Local S
Matches 20
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15-MAR-2000;
23-JUN-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The sequence represents a sequencing primer used to sequence an exon of the human CYP2B6 gene. It is used for specific detection and genotyping of CYP2B6 alleles in humans, determination of which is useful for the optimisation of therapies utilising CYP2B6 substrates. Oligonucleotide sequences are useful in detection of the individual predisposition to several common cancers caused by environmental carcinogens, and diseases treated with drugs that are targets of the CYP2B6 gene product, whose metabolism is therefore dependent on CYP2B6. Cancer or susceptibility to cancer can be diagnosed by detecting the presence of a molecular variant of CYP2B6. From variants of the alleles, modulators of the activity can be developed for use in treatment and prevention of CYP2B6-related
                                                                         Treating level, a
                                                                                                                                                                                                                                                                                                                                                            High
                                                 transcriptional activity.
                                                                                                                                                                                                                                                         01-SEP-2000;
                                                                                                                                                                                                                                                                                   08-MAR-2001
                                                                                                                                                                                                                                                                                                           WO200115676-A2
                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                    Polymorphic sequence for ABC1 polymorphic site #38.
                                                                                                                                                                                                                                                                                                                                                                                                                17-MAY-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                         AAF93028;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAF93028 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      polynucleotide(s) of the polymorphic human CYP2B6 gene for sction and treatment of disorders i.e. cancer.
                                                                                                                                                                                                                                                                                                                                                            density
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20; Conserv
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                                                            a lower than normal high density lipoprotein-cholesterol (HDL-C) higher than normal triglyceride level, or a cardiovascular by administering a compound that modulates LXR- or RXR-mediated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GATTACAGGCATGAGCCACCA 21
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ilarity 95.2%;
Conservative
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                                                                                                                                         Brooks-Wilson AR,
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2000US-00526193.
2000US-0213958P.
                                                                                                                                                                                                                                                         2000WO-IB001492.
                                                                                                                                                                                                                                                                                                                                                            lipoprotein-cholesterol; HDL-C; cardiovascular;
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Pred. No. 1.
                                                                                                                                         Pimstone SN,
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Disclosure;

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317pp; English.

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CCAAAGTGCTGGGATTACAGG 407

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Query Match Best Local S Matches 20

Similarity 20; Conser

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Score 19.4; Pred. No. 1

.3e+03

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RESULT 61
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                                                       The invention relates to a method for the amplification of DNA that maintains genes and copy number of the sequence. This method is useful for easy and operable amplification of DNA. The method was carried out by fragmentation genomic DNA, preparation of blunt end of the fragmented DNA, ligation of an adapter to the blunted DNA, PCR of the ligated DNA in 2 steps, and confirmation of the amplified APC gene. The current sequence represents a PCR primer used in an example from the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (TAKA-)
(KOKU-)
(IYAK-)
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  Sequence
                                                                                                                                                                                                                                                                                                                                         Amplification of DNA maintaining genes and copy number of the a genome, and their ratios in the resultant DNA fragment.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
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20; Conserv
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KOKURITSU GAN CENT SOCHO.
IYAKUHIN FUKUSAYO HIGAI KYUSAI KENKYU SH.
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     22
                                                                                                                                                                                                                                                                                    SEQ ID
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  B₽;
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  Α.
                                                                                                                                                                                                                                                                                    NO 50; 33pp; Japanese
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  4 C; 7 G; 4 T;
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Pred. No. 1
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  0 U;
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     0 Other;
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RESULT 621
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AAQ73577
ID AAQ735
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AAQ73576
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Matches 20
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25-JUN-1995
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                                                                                                                                            This enhancer element is part of a DNA construct used for treating human carcinoma which contains a cancer therapeutic protein under the control of a promoter and 3 enhancer sequences in a specific 5'-3' order. This enhancer element is derived from the flanking region of the human epithelial cell cytokeratin-8 gene. (Updated on 25-MAR-2003 to correct Pi
                                                                                                                                                                                                                                                                                                                                                                 misc_difference
                                                                                                                                                                                                                                                                                                                                                                                                                       Enhancer element; carcinoma; secretory leukoprotease-inhil
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                                                                                                                                                                                                                                                                                                                                                                                                          Homo
                                                                                                                          Sequence
                                                                                                                                                                                                          DNA construct for treating human carcinoma - includes a cancer-therapeutic gene under the control of a promoter and a gp. of enhancer
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25-MAR-2003
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                                   standard; DNA;
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  entry)
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                                                                                                      Score 19.4; DB 1;
Pred. No. 1.3e+03;
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                                                                                                                           3 Other;
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RESULT 622
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 misc_difference
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Synthetic
                           S182 gene; familial Alzheimer's
polymerase chain reaction; PCR;
                                                                      Primer Alu 5'
                                                                                                  17-JUN-1997
                                                                                                                           AAT63214;
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                                                                                                                                                                                                                                                                                                    2.0%;
                                                                                                                                                                                                                                                                                                                                                8 C; 6 G; 4 T; 0 U; 3 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           purine
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           human carcinoma -
control of a promo
                                                                                                                                                            ВÞ
                                                                                                                                                                                                                                                                                  Pred. No. 1.30
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  tumor; cancer;
                                                                                                                                                                                                                                                                                                         Score 19.4;
Pred. No. 1
                            disease; diagnosis; transgenic primer; artificial chromosome;
                                                                        PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   gene; cytokeratin gene-8; ss.
                                                                                                                                                                                                                                  24
                                                                         for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            promoter and
                                                                        PAC isolation.
                                                                                                                                                                                                                                                                                                         .3e+03;
                                                                                                                                                                                                                                                                                                                        DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   gene;
                                                                                                                                                                                                                                                                                                                      Length
                                                                                                                                                                                                                                                                                           Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              gp.
                                                                                                                                                                                                                                                                                                                        24;
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                               enimal;
PAC; ss.
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WO9703999-A1

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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Inter-Alu PCR was performed on YACs 905C2 and 763B11. Unpurified YAC DNA was amplified with generate primers Alu 5' (AAT63214) and Alu 3' (AAT63215). Genetic linkage strategies have placed a gene causing early onset Alzheimer's disease (AD) on the long arm of chromosome 14 between D14S289 and D14S61. The gene, S182 (see also AAT63207), was localised to a 100 kb region between D14S77 and D14S668E (see also AAT63216-22). A number of novel mutations in the S182 gene have been identified in
Detection of single nucleotide polymorphisms in genomes by preparation
                        WPI; 2000-293181/25
                                                                                                                                                                                                                        Human; single nucleotide polymorphism; SNP; genotyping; DNA analysis; allele specific oligonucleotide; ASO; reduced complexity genome; RCG; genomic classification; identification; DNA fingerprinting; tumour characterisation; hybridisation; ss.
                                                                                                                                                                                                                                                                                                                                        AAA35956;
                                                                                                                                                                                                                                                                                                                                                               AAA35956 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 5 A; 4 C; 6 G; 3 T; 0 U; 2 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 2; Page 11; 26pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New mutants of the S182 gene associated with familial Alzheimer's disease - and related protein and transgenic animals, useful as models for screening and assessing potential drugs.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1997-132571/12.
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02-AUG-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                26-JUN-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         06-FEB-1997.
                                                                                                                         24-SEP-1999;
                                                                                                                                                                                                                                                                                      Human genomic
                                                                                                                                                                                                                                                                                                                26-JUL-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     families
                                                                                                 25-SEP-1998;
                                                                                                                                                                          WO200018960-A2
                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (UNIW ) UNIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                   868
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            18;
                                                                          MASSACHUSETTS INST TECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   GGATTACAGGCGTGAGCCAC 887
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   multiply affected by early onset AD
                                                                                                                                                                                                                                                                                                                                                                                                                             GGATTACAGGYRTGAGCCAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hardy JA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WASHINGTON SCHOOL MED SOUTH FLORIDA.
                                                                                                                                                                                                                                                                                                              (first entry)
                                                Jordan
                                                                                                                                                                                                                                                                                       SNP allele specific oligonucleotide SEQ ID NO:13.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             95US-0001500P.
                                                                                                 98US-0101757P
                                                                                                                          99WO-US022283
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                                                                                                                                                                                                                                                                                                                                                               DNA;
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                                                œ
                                                  Housman
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                                                                                                                                                                                                                                                                                                                                                               ВР
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Pred. No. 1.
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                                                  Charest
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RESULT 624
AAA27180/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A method has been developed for detecting the presence or absence of a single nuclectide polymorphism (SNP) allele in a genomic sample. The method comprises preparing a reduced complexity genome (RCG) from the genomic sample and analysing the RCG for the presence or absence of a SNP allele. The method can be used to characterise a tumour, to generate a genomic pattern for an individual genome or to generate a genomic classification code for a genome. The method can be used to assess whether a subject is at risk for developing a disease or to identify a set of SNP alleles associated with a disease. The method can also be used to perform linkage analysis. AAA35944 to AAA35947 represent sequences used in the exemplification of the present invention. AAA35948 to
A novel method for simultaneously determining the level of a number of target polynucleotides in a sample has been disclosed. The method involves forming double stranded copies of the target sequence in direct proportion to the target levels in the original sample. The target sequence is copied using primer pairs designed to flank a defined region in the target sequence. The double stranded copies are then cleaved and reacted with either first or second adaptor sequences. The first and
                                                                                                                                                                               Measuring target polynucleotide sequences in biological samples by contacting sequence-selective primer pairs, forming conjugates with adaptor molecules, polymerizing target-identifier dimers and quanti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                P2; CX5C chemokine; Chromosome 5q31; gene therapy; asthma; PCR primer; allegic rhinitis; urticaria; anaphylactic shock; hives; hay fever; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAA27180;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAA36632 represent nucleotide sequences containing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     and analysis of reduced complexity genomes, useful for genotyping, fingerprinting and determining allele frequency of SNPs.
                                                                                                                                    Disclosure;
                                                                                                                                                                                                                                                                                        Dolganov
                                                                                                                                                                                                                                                                                                                                                             16-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                              12-NOV-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Forward primer P2 for target
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                                                                                                                                                                                                                                                         2000-387825/33.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      standard; DNA;
                                                                                                                                 Page 99; 103pp; English.
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                                                                                                                                                                                                                                                                                         Novikov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                             98US-00193320.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sequence human P2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 19.2;
Pred. No. 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     gene.
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                                                                                                                                                                                    quantifying
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RESULT 625
AAH4560/C
ID AAH456
XX AAH456
XX AAH456
XX AAH456
XX Protes
XX W02001
XX W02001
XX U7FU
PA (SHAN
XX YX
XX PI MAO Y
XX WPI;
XX Human
PT inmunn
PT inmunn
PT infla
XX Human
PT infla
XX Examp
XX This
CC CDNA
CC This
CC CDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cc through the target sequences. The adaptor sequences are then polymerised to conjugate mixtures are then allowed to form dimers with each other through the target sequences. These dimers are then polymerised to form cd dimer multimers. The relative abundances of target identifiers in the continuous polymucleotide abundance level profiles for cells and contisues under various conditions, stages of development and and disease consistency particularly where the target polymucleotide is present at low levels. The method may also be used in the discovery and evaluation of condition to the method described above, the invention also includes the colymucleotide and polypeptide of P2. P2 is thought to be a member of a convel chemokine family, denoted CXSC and may be associated with immune convel chemokine family, denoted CXSC and may be associated with immune convel chemokine family, denoted CXSC and may be associated with in the convel chemokine family, denoted CXSC and may be useful in the convel chemokine family, denoted CXSC and may be associated with immune convel chemokine family, denoted CXSC and may be useful in the convel chemokine family, denoted CXSC and may be useful in the convel chemokine family, denoted CXSC and may be useful in the convel chemokine family, denoted CXSC and may be useful in the converse of the pay. The pay polymucleotide may be useful in the converse of the pay. The pay polymucleotide to chromosome 5, within the cytokine gene cluster at 5331. The present sequence is the forward primer pay for target sequence human P2 gene has been localised to chromosome 5, within the cytokine gene cluster at 5331. The present sequence is the forward primer pay.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Simi
Matches 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Protease regulatory protein 9; malignant tumour; had HIV infection; immunological disease; inflammation; virucide; immunomodulatory; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                     Human protease regulatory protein 9 and encoded polynucleotide, diagnosis and treatment of malignant tumors, hemopathy, human immunodeficiency virus infection, immunological diseases and
                                                                                                                                                                                                                                                                                                                                                    мао Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       29-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-DEC-2000; 2000WO-CN000652
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12-JUL-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PCR primer specific for human protease regulatory protein 9 cDNA
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                                                                                                                                                                                                                                                                                                                                                                                                      (UYFU-)
                                                                                                                                                                                                                                                                                                  WPI; 2001-441850/47
                                                                                                                                                                    inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         676 CACTGCAACCTCTGCCTCCCGGGT 699
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                                                                                                                                                                                                                                                                                                                                                                                                         SHANGHAI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 4 A; 3 C; 12 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       99CN-00127227
                                                                                                                                                                                                                                                                                                                                                                                                         BIO DOOR GENE TECHNOLOGY LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1.9%;
87.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 19.2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       .4e+03
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     haemopathy;
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Example 3; Page

17;

35pp;

Chinese

invention relates to human protease regulatory protein 9, an encoding it. The invention includes a vector containing the cell transformed with the vector, and an antibody directed a

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RESULT 626
AAH46154/c
ID AAH461
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This sequence represents cDNA encoding cysteine protease 10. The protein control of 10 kp, and has homology with a cysteine protease given in AAB73746 over a 51 amino acid stretch. The invention crelates to cysteine protease 10 (AAB73746), nucleic acids encoding it (AAH46153), and a method for the recombinant production of cysteine protease 10. The present invention additionally discloses an agonist of cysteine protease 10 for therapeutic use, and an antibody which specifically binds to cysteine protease 10. Cysteine protease 10, and conclectides which encode it may be used for treating a variety of conditions. The protein may also be used to screen for modulators of its conditions. The protein may also be used to screen for modulators of its cativity or for peptide fingerprinting identification. The polynucleotide can be used as a primer for nucleic acid amplification reactions or as a probe for hybridisation reactions, or in producing gene chips or conditions. Sequences AAH46154-AAH46155 represent reverse transcription-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 the protein. The protein and polynucleotide sequences can be used in t diagnosis and treatment of malignant tumours, haemopathy, human immunodeficiency virus (HIV) infection, immunological diseases, and various inflammatory conditions. Use of the protein or polynucleotide
                                                                                                                                                                                                                                                                                                                        Cysteine protease 10 and encoded polynucleotide, applicable in diagnosis and treatment of malignant tumor, hemopathy, HIV infection, immunological diseases and various inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cysteine protease 10; human; recombinant production; malignant tumour; cancer; blood disease; HIV infection; human immunodeficiency virus;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            immune disorder; inflammatory condition; cytostatic; anti-HIV;
antiinflammatory; immunomodulator; reverse transcription-PCR;
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                                                                                                                The invention relates to the human Pax protein 9 with cytostatic, virucide, immunomodulatory, antiinflammatory, haemostatic and anti-HIV activity. The polypeptide and encoded polynucleotide, with paired box domain, are applicable in diagnosis and treatment of malignant neoplasm, haemopathy, human immunodeficiency virus (HIV) infection, immunological diseases, various inflammations, developmental disorders, growth developmental disturbance and Waardenburger's syndrome. The polynucleotide is useful for gene therapy. The present sequence is that of a human Pax protein 9 PCR primer, useful to the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; Pax protein 9; cytostatic; virucide; immunomodulatory; anti-HIV; paired box domain; neoplasm human immunodeficiency virus; HIV; infection; immunological disease; developmental disorder; growth developmental disturbance; Waardenburger's syndrome; gene therapy; PCR primer; ss.
                                                                                             Sequence
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                                                                                                                                                                                                                                                                New human Pax protein 9 for diagnosing and treating developmental disorders, malignant neoplasm, hemopathy, human immunodeficiency infection, immunological diseases and various inflammations.
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15-MAR-2000;
23-JUN-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to a method for treating a patient diagnosed as having a lower than normal high density lipoprotein-cholesterol (HDL-C) level, a higher than normal triglyceride level, or a cardiovascular disease, involving administering a compound that modulates LXR- or RXR-mediated transcriptional activity or ABCl expression or activity. The LXR gene product may be used in an assay to identify compounds useful for the treatment of a disease or condition selected a lower than normal HDL cholesterol level, a higher than normal
                       Pumarase 9; cytostatic; antiviral; immunomodulatory; antiinflammatory;
cardiant; cancer; haemopathy; human immunodeficiency virus; HIV;
infection; immunological disease; inflammatory disease; PCR primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Treating a lower than normal high density lipoprotein-cholesterol (HDL-C) level, a higher than normal triglyceride level, or a cardiovascular disease, by administering a compound that modulates LXR- or RXR-mediated
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Pred. No. 1
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primer; ss

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RESULT 6
AAF27674
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to an isolated polypeptide of Fumarase 9 comprising the 80 amino acid sequence defined in the specification, or its fragment, analogue or derivative. The polypeptide and the polymucleotide encoding it are useful in the diagnosis and treatment of malignant neoplasms, haemopathy, HIV infection, immunological diseases and various inflammatory diseases. The present sequence is a primer used to isolate a polynucleotide encoding the polypeptide of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mao
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Fumarase 9 polynucleotide and polypeptide, useful in diagnosis and treatment of malignant neoplasm, hemopathy, HIV infection, immunoldiseases and various inflammatory diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unidentified
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200148176-A1
        Determining whether a subject has or is predisposed to disease with IL-1 polymorphism involves determining presence of marker comprising IL-1 inflammatory haplotype.
                                                                                                                                                                                                                               IL-1; interleukin; inflammation; infection; ss
                                                                                                                                                                                                                                                      Primer
                                                                                                                                                                                                                                                                            02-APR-2001
                                                                                                                                                                                                                                                                                                  AAF27674;
                                                                                                                                                                                                                                                                                                                     AAF27674
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 3; Page 11; 34pp; Chinese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18-DEC-2000; 2000WO-CN000612
                                                                                                                                          30-JUN-2000; 2000WO-US018318
                                                                                                                                                                                     WO200100880-A2
                                                                                                                                                                                                          Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (BIOW-) BIOWINDOW GENE DEV LTD
                                                                                                                     30-JUN-1999;
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                                                                                               INTERLEUKIN GENETICS
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                                                                                                                                                                                                                                                                                                                                                                                       TTTTTAATTTTTGAGACAGAGTC 630
                                                                                                                                                                                                                                                                                                                      standard;
                                                                                                                                                                                                                                                                                                                                                                             TTTTTGGTTTTTTGAGACGGAGTC
                                                                         Cox
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP; 12 A; 7 C; 2 G; 3 T; 0 U; 0 Other;
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                                                                                                                     99US-00345217
                                                                         Camp
                                                                                                                                                                                                                                                                                                                      DNA;
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                                                                            Di Giovine
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Pred. No. 1
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                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 24;
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RESULT 631
AAH40034/c
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to a new method for determining whether a subject has or is predisposed to developing a disease or condition that is associated with an IL (interleukin)-1 inflammatory haplotype, comprises detecting at least one allele of the haplotype, where the presence of the allele indicates that the subject is predisposed to the development or has the disease or condition. The method is useful for determining whether a subject has or is predisposed to inflammatory disease, a degenerative disease, an immunological disorder, an infectious disease, trauma induced disease, or cancer. The above conditions associated with an IL-1 inflammatory haplotype can be treated or prevented by administering a therapeutic that compensates for a causative mutation that is in linkage disequilibrium with at least one II-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                polycystic kidney disease; osteogenesis imperfecta; autoimmune disease;
acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ANS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 24 BP; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAH40034;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAH40034 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         polymorphism
                                                                                                          Claim 1; Page 64; 83pp; English.
                                                                                                                                               absence or identity of single polynucleotide polymorphism
                                                                                                                                                              New genotyping oligonucleotide, useful for detecting the presence,
                                                                                                                                                                                         WPI;
                                                                                                                                                                                                                   Picoult-Newburg
                                                                                                                                                                                                                                                                          15-OCT-1999;
                                                                                                                                                                                                                                                                                                    13-OCT-2000; 2000WO-US028436
                                                                                                                                                                                                                                                                                                                                 26-APR-2001.
                                                                                                                                                                                                                                                                                                                                                          WO200129262-A2
                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                inflammation; forensic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          868 GGATTACAGGCGTGAGCCACCACG 891
                                                                                                                                   sample.
                                                                                                                                                                                         2001-290930/30
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                                                                                                                                                                                                                                               ORCHID BIOSCIENCES INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GGGATACAGGCGTGAGCCACCGCG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               lower PCR primer SEQ ID 2830.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                          99US-0160096P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 7 C;
                                                                                                                                                                                                                                                                                                                                                                                                                 investigation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24
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); Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19.2;
No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                 paternity analysis; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        24;
                                                                                                                                                  in a nucleic
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primer extension (SNPE) primers, and the sequences of regions flanking sites of single nucleotide polymorphisms SNPs. The present invention includes kits for determining the presence or absence of a SNP, using the oligonucleotides of the invention. The PCR primers are used to amplify a SNP flanking sequence, the SNPE primer is used as a genotyping primer. The oligonucleotides are useful for genotyping a nucleic acid sample by

Sequences AAH37205 -

AAH40944 represent PCR primers, single nucleotide

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Query Match
Best Local
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New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nuc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; lesch Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis; inflammation; forensic investigation; paternity analysis; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SNP specific upper PCR primer. SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAH37609 standard;
                                                                                                                                                    Picoult-Newburg L,
                                                                                                                                                                                                                                                                                     15-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Single nucleotide polymorphism; SNP; single nucleotide primer extension;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14-AUG-2001
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                                                                                                                                                                                                                                                                                                                                               13-OCT-2000; 2000WO-US028436
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200129262-A2
                                                                                                                                                                                                                           (ORCH-) ORCHID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         a human SNP containing DNA sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       158
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP; 6 A;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first
                                                                                                                                                                                                                           BIOSCIENCES INC
                                                                                                                                                                                                                                                                                     99US-0160096P
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                                                                                                                                                             Pohl M;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            24
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Pred. No. 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1;
polymorphism in a nucleic
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performing a single-nucleotide primer extension reaction. The oligonucleotides are useful for determining the presence, absence or identity of a SNP and for genotyping nucleic acid samples, for e.g. to assess by association analysis the genotype of an individual or group of individuals, having a pathological phenotypic trait suspected of being caused by one or more SNPs. Phenotypic traits include diseases e.g. agammaglobulinaemia, diabetes insipidus, Lesch-Nyhan syndrome, muscular dystrophy, familial hypercholesterolaemia, polycystic kidney disease, osteogenesis imperfecta and acute intermittent porphyria. Phenotypic raits also include symptoms of or suscensivibility to multifearchis!
                                                                                                                                       traits also include symptoms of or susceptibility to multifactorial disease of which a component is or may be genetic such as autoimmune diseases, including, rheumatoid arthritis, multiple sclerosis, inflammation, cancer, nervous system diseases and infection by pathogenic microorganism. The method is also useful in forensic investigations and paternity analysis. The present sequence represents a PCR primer specific
Sequence 24 BP; 4 A; 8 C; 3 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The oligonucleotides are useful for genotyping a nucleic
                                                                                                           human SNP
                                                                                                containing
                                                                                                     DNA sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sample by
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Query Match
Best Local :
                                  Matches
                 670
Н
                                           Similarity
                 TTGGCTCACTGCAACCTCTGCCTC 693
                                 Conservative
                                         1.9%;
                                          Score 19.2;
Pred. No. 1
                                  Mismatches
                                           1.4e+03
                                                   DB 1;
                                                  Length 24;
                                  0
                                  Gaps
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RESULT 633 AAF81133/c single nucleo inflammation; Human; prostaglandin-endoperoxide synthase 2; PTGS2; cyclooxygenase AAF81133 standard; DNA; 24 Homo sapiens. PCR primer used 02-MAY-2001 nucleotide polymorphism; SNP; (first entry) PCR 6 primer; ss amplify PTGS2 exon immune-related φ SEQ Ħ disorder; arthritis;

WO200107662-A1

99US-0145170P.

24-JUL-2000; 2000WO-US020114.

(GENA-) GENAISSANCE PHARM INC

RR, Nandabalan **7** Sanchis Þ Stephens Ĺ,

New nucleic acid containing polymorphisms for gene therapy of inflammation and for ε s in the cyclooxygenase-2 gene, establishing a genotype or

Example 1b; Page 39; 118pp; English.

AAF80897 represent human PTGS2 gene and coding sequence, and the proprotein is represented by AAB72199. The invention includes PCR and sequencing primers, and probes represented in AAF80898 - AAF81151 v are used to isolated and characterise the PTGS2 gene sequence, and This invention relates to a polynucleotide sequence that is a polymorphic variant of the human prostaglandin-endoperoxide synthase 2 (PTGS2) gene also referred to as cyclooxygenase 2. The human PTGS2 gene sequence AAF800896 contains 27 single nucleotide polymorphisms (SNPs). AAF80896 and AAF81151 which

Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide primer extension (SNPE) primers, and the sequences of regions flanking sites of single nucleotide polymorphisms SNPs. The present invention includes kits for determining the presence or absence of a SNP, using the oligonucleotides of the invention. The PCR primers are used to amplify a

flanking sequence,

SNPE

primer

Claim 1;

Page 52; 83pp;

English.

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RESULT 634
AAI64727/c
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Best Local S
Matches 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             locate the positions of the SNPs. PTGS2 proteins and polynucleotide sequences are used to express variant PTGS2 proteins, for structural analysis or drug-binding studies and also in gene therapy (either expressing PTGS2 or inhibitory RNA). Antibodies raised against PTGS2 are useful for diagnosis, prognosis and therapy and analysis of the new, and known, polymorphisms and used to determine PTGS2 haplotype and genotype, especially for determining association between a particular trait, e.g. a clinical response to drugs that target PTGS2 but also disease susceptibility, severity or stage. Anti-PTGS2 antibodies are particularly used for developing diagnostic tests and treatments for immune-related disorders such as archritis and inflammation. The polymorphisms may also be used to study expression and biological function of PTGS2. Transgenic animals that express PTGS2 are used to study expression of PTGS2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; line 1-12; cytostatic; virucidal; immunomodulatory; antiinflammatory; haemostatic; malignant tumour; HIV; infection; human immunodeficiency virus; immunological disease; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 07-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAI64727;
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                               The invention relates to human line 1-12 with cytostatic, virucidal, immunomodulatory, antiinflammatory and haemostatic activity. The protein and encoding polynucleotide are used in diagnosis and treatment of malignant tumour, haemopathy, human immunodeficiency virus (HIV) infection, immunological diseases and various inflammations. The present sequence is that of a human line 1-12 PCR primer, useful to the invention
                                                                                                                                                                                            Line 1-12 and encoded polynucleotide, used in diagnosis and treatment malignant tumors, hemopathy, human immunodeficiency virus infection,
                                                                                                                                                                                                                                            WPI; 2001-597126/67.
                                                                                                                                                                                                                                                                             Mao Y,
                                                                                                                                                                                                                                                                                                                                             27-MAR-2000; 2000CN-00115143
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                                                                                                                                                                           immunological diseases
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                                                                                                                                            Example
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                                                                                                                                                                                                                                                                             Xie Y;
                                                                                                                                                                                                                                                                                                              SHANGHAI BIOWINDOW GENE DEV INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      for in vivo drug screening and testing,
                                                                                                                                          Page 16; 33pp; Chinese.
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                                                                                                                                                               hemopathy, numer.

----- and inflammation.
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Pred. No. 1
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Query Match

Sequence

24

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4 C; 1.9%;

10

<u>ი</u> 5 T;

0 U;

Score 19.2;

DB 1; 0 Other;

Length

24;

Human; sailor transposase 9.35; recombinant cancer; tumour; HIV infection; human immunoc

nbinant production; gene immunodeficiency virus;

therapy;
cytostatic;

invention

Human

sailor

transposase

9.35 RT-PCR primer,

SEQ ID NO:3.

of.

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RESULT 635
ABQ83629
ID ABQ836
XX ABQ836
XX ABQ836
XX ABQ836
XX ABQ836
XX ABQ836
XX Human
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XX Human
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XX PCR p1
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XX Homo E
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PN CN1345
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                                                                                                                                                                                                                        RESULT 636
ABV76754
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                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention describes human mPer3-10.01 (I). Also described a method for producing (I) using DNA recombination technology. (I) car used in the treatment of several diseases, such as vegetative nervous dysfunction, psychic disease, endocrinopathy, growth development disturbance disease and tumours. The present sequence represents a PCF disturbance disease and tumours.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; mPer3-10.01; vegetative nervous dysfunction; psychic disease; endocrinopathy; growth development disturbance disease; tumour; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel polypeptide-human mPer 3-10.01 polypeptide.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    primer
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                                                                                                                 28-MAR-2003
                                                                                                                                                          ABV76754
                                                                                                                                                                                                   ABV76754 standard; DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP; 4 A; 8 C; 6 G; 6 T; 0 U; 0 Other;
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Pred. No. 1
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RESULT 637
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Matches 21
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                                                 Mao
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WPI; 2002-637132/69
                                                                                                                                                                                                                                                                                                 CN1352077-A
                                                                                                                                                                                                                                                                                                                                                                                              Human; transcription regulatory factor 16.06; tumour; embryonic development malformation; protein metabolic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human transcription
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                                                                                               BODE GENE DEV
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disorder disease.
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Matches 21
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                                                                                                 The invention relates to the novel human alpha 2,3-sialyltransferase 9.90, and the polymucleotide encoding it. The polymeptide is useful for treating various diseases, such as an immunological defect, various tumours and various inflammations. The invention also discloses the antagonist resisting the polymeptide and its treatment effect. The present sequence represents a PCR primer used to amplify the the human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention describes human transcription regulatory factor 16.06 (I). Also described is a DNA recombination process used to produce (I). (I) can be used for treating various diseases, such as embryonic development malformation, tumours and protein metabolic disorder disease. The present sequence represents a PCR primer for (I), which is used in an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New polypeptide-human transcription regulatory factor 16.06 and polynucleotide for encoding the polypeptide, embryonic development malformation, tumors, and protein metabolic disorder disease.
                                                       Sequence
                                                                                 present sequence represents a PCR primer used to amplialpha 2,3-sialyltransferase 9.90 gene of the invention
                                                                                                                                                                                                                                New polypeptide-human alpha 2,3-sialyltransferase 9.90 for treating immunological defect, various tumors and various inflammations.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example
                                                                                                                                                                                                     Example 2; Page 19 (Disclosure); 33pp; Chinese.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; alpha 2,3-sialyltransferase 9.90; immunological defect; tumour;
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                                                                                                                                                                                                                                                                                                        Xie Y;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              standard;
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             1.9%;
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Pred. No. 1.4e
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                                                         G, 8 T; 0 U; 0 Other;
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Pred. No. 1.4e+03;
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention provides the protein and coding sequences of human 6 grotein subunit 9.02. The sequences can be used in the treatment of cancers, coughs, cardiac asthma, diarrhoea, constipation, colic, psychic disease and morphinic analgesic acute poisoning. The present sequence is a PCR primer used to isolate the coding sequence of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CN1345751-A.
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morphinic analgesic acute poisoning; ss.
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                                                                                                     ss; guanosine triphosphatase activator protein 10.12; PCR; primer; malignant tumour; inflammation; immunological disease; haemopathy; human immunodeficiency virus infection; HIV; RT-PCR;
                                                                                                                                                                                                                  Guanosine
                                                                                                                                                                                                                                                                   04-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                        ABX14631 standard; DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10-NOV-2000; 2000CN-00127330.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     recombinant production; gene therapy; can human immunodeficiency virus; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ras GTP enzyme-activating protein 20.68; cancer suppressor protein-20.68; recombinant production; gene therapy; cancer; tumour; HIV infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        07-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 24 BP; 5 A; 5 C; 10 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                      Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                             RT-PCR; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ras GTP enzyme-activating protein 20.68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABV76761;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABV76761 standard;
                                                      мао Y,
                                                                                                                                                                   05-JAN-2001; 2001CN-00105082
                                                                                                                                                                                                                             05-JAN-2001; 2001CN-00105082
                                                                                                                                                                                                                                                                                      14-AUG-2002.
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                                                                                                                CO LTD SHANGHAI
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Pred. No. 1.
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WPI; 2002-742044/81

24;

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RESULT 642
ABV99633
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to Ras GTP enzyme-activating protein 20.68 (ABPSB53) and nucleic acids encoding it (ABPN6760). The protein has a molecular weight of 20.68 kD and is also referred to as cancer suppressor protein-20.68. The invention also relates to a method for the recombinant production of the protein, an antagonist of the protein, and the use of the protein, gene and antagonist in therapeutic applications. Ras GTP enzyme-activating protein 20.68 can be used in the treatment of a variety of diseases such as cancer and HIV (human immunodeficiency virus) infection. Sequences ABV76761-ABV76762 represent reverse transcription-PCR (RT-PCR) primers used in an exemplification of the invention to isolate Ras GTP enzyme-activating protein 20.68 cDNA
Sequence
                         peptide receptor 11.66 PCR
                                     The invention relates to human natriuretic peptide receptor 11.66. The polypeptide is useful for treating various diseases, such as malignant tumours, haemopathy, HIV infection, immunological diseases and various inflammations. The present sequence is that of a human natriuretic
                                                                                                                                                        New polypeptide-human natriuretic peptide 11.66 and polynucleotide encoding the polypeptide, useful for treating malignant tumors, hemopathy, HIV infection, immunological diseases and various
                                                                                                                                                                                                                   WPI; 2002-644453/70.
                                                                                                                                                                                                                                                Мао У,
                                                                                                                                                                                                                                                                                                                                                                 05-JUN-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; natriuretic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human natriuretic peptide receptor 11.66 PCR primer
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                                                                                                              Example 2; Page 16 (Disclosure); 31pp;
                                                                                                                                                                                                                                                                                                        02-NOV-2000; 2000CN-00127189
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                                                                                                                                              inflammations.
                                                                                                                                                                                                                                                                            (BODE-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               n; natriuretic peptide receptor 11.66; receptor; tumour; haemopathy; human immunodeficiency virus; infection; immunological disease;
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                                                                                                                                                                                                                                                Xie Y;
                                                                                                                                                                                                                                                                            BODE GENE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GCGATTCTCCTGTCTCAGCCTCCC 1027
24 BP; 3 A; 7 C; 7 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GTGATTCTCCTACCTCAGCCTCCC 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      PCR;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    primer; ss.
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                           primer useful
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Pred. No. 1.
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                                                                                                                 Chinese.
                            'n
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                           examples of the invention
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RESULT 644
AB199962
ID AB1999
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AC AB1999
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ABI99962 standard;

DNA;

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ABI99962;

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RESULT 643
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Best Local S
Matches 21
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Best Local Similarity
Matches 21; Conserv
                                                                                                                        The invention relates to an isolated polypeptide of phytochrome 12, the cDNA encoding it, and its fragment, analogue or derivative. Also included are vectors expressing protein, a host cell comprising the vector, the isolation of modulators of the protein and an anti-phytochrome 12 antibody. The protein and nucleic acid are used in diagnosis and treatment of a malignant tumour, haemopathy, human immunodeficiency virus (HIV) infection, immunological diseases and various inflammations. The present sequence is an RT-PCR (reverse transcriptase PCR) primer used to isolate the cDNA encoding the phytochrome 12
                                                                                                                                                                                                                                                                                 phytochrome 12 and encoding polynucleotide, used in diagnosis and treatment of malignant tumors, hemopathy, human immunodeficiency virus infection, immunological diseases and inflammation.
                                                                                                                                                                                                                                                                                                                                                               Mao
                                                                                                   Sequence
                                                                                                                                                                                                                                                        Example 2; Page 17; 36pp; Chinese.
                                                                                                                                                                                                                                                                                                                                    WPI; 2002-083184/11.
                                                                                                                                                                                                                                                                                                                                                                                                                 24-MAY-2000; 2000CN-00115823.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; ss; phytochrome 12; malignant tumour; haemopathy; human immunodeficiency virus infection; HIV; immunological
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        09-APR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAS20139;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200192316-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               inflammations;
                                                                                                                                                                                                                                                                                                                                                                                       (SHAN-) SHANGHAI BIOWINDOW GENE
                        1001 CAAGCGATTCTCCTGTCTCAGCCT 1024
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               phytochrome
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                                                             Similarity
                                                                                                   24 BP; 4
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CAAGCGATTCTTGTGCCTCAGCCT
                                                  Conservative
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                                                                                                   A; 8 C; 5
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                                                            1.9%;
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                                                                                                   G; 7 T;
                                                Score 19.2; I
Pred. No. 1.46
0; Mismatches
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Pred. No. 1.4e+03;
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                                                                                                    0 Other;
                                                               4e+03;
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RESULT 645
AA172742
ID AA17277
XX AA1727
XX AA1727
XX O3-JUL
XX Gene,
KW Human
XX Gene,
KW Human
XX Gene,
XX OGNO
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XX WO2001
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XX WO3-APP
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to human phosphatidic acid phosphatase 2-12 (AAM49149) and to nucleic acids encoding it (ABI99960). The protein has a molecular weight of 12 kD. The inevntion also relates to a method for the recombinant production of the protein, an antagonist of the protein, and the use of the protein, gene and antagonist in therapeutic applications. Phosphatidic acid phosphatase 2-12 can be used in the treatment of a variety of diseases such as cancer and HIV (human immunodeficiency virus) infection. Sequences ABI99961-ABI99962 represent reverse transcription-PCR (RT-PCR) primers used in an exemplification of the invention to isolate human phosphatidic acid phosphatase 2-12 cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New polypeptide-human phosphatidic acid phosphotase 2-12 diseases such as cancer and human immunodeficiency virus
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                                                                                                                             primer; ss.
                                                                                                                                          Gene; human cytokine receptor 15; malignant tumour; haemopathy; human immunodeficiency virus; HIV; inflammation; gene therapy;
                                                                                                                                                                                                                03-JUL-2002
                                                                                                                                                                                                                                                                       AAI72742 standard; DNA; 24 BP
              23-APR-2001; 2001WO-CN000577
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                                            08-NOV-2001
                                                                       WO200183536-A1
                                                                                                  sapiens.
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                                                                                                                                                                                     cytokine
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                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
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                                                                                                                                                                                                                                                                                                                                               TTTTTAGTAGAGACAGGGTTTCAC
                                                                                                                                                                                                                                                                                                                                                                           TTTGTGGTAGAAACAGGGTTTCAC 333
                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP; 6 A;
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                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                    receptor 15 PCR primer #1.
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                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                      No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                   1.4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         virus; gene therapy;
CR; RT-PCR; primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                  Length
                                                                                                                                            gene therapy; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                    24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  for treating infection.
                                                                                                                                                                                                                                                                                                                                                                                                       0;
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RESULT 646
AAL43822
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Best Local S
Matches 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention relates to human cytokine receptor 15 (see AAB47984). Human cytokine receptor 15, and the DNA encoding it, are in diagnosis and treatment of mallignant tumour, hemopathy, human immunodeficiency virus (HIV) infection, immunological diseases and various inflammations. This sequence is a PCR primer which was used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cytokine receptor 15 and encoded polynucleotide, applicable in and treatment of developmental disorders, cancer, hemopathy, HI infection, immunological diseases and various inflammations.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2002-026253/03
                                                                                                                                                                                                                                                                                                                                                              Human, ss, gene therapy; oncogene protein 11.66; malignant tumour; haemopathy; development disturbance; HIV; immunological disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 3; Page 12; 38pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       27-APR-2000; 2000CN-00115491.
                    The invention comprises the amino acid and coding sequence of toncogene protein 11.66. The oncogene protein 11.66 DNA and protogenences are useful for treating malignant tumour, haemopathy, development disturbance, HIV infection, immunological disease a development disturbance.
                                                                                                                                                                                                                                                                         30-JAN-2002.
                                                                                                                                                                                                                                                                                                                                                                                                Human oncogene protein 11-66 PCR primer 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAL43822;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAL43822 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                example
protein 11.66 PCR primer
                                                                                                          Novel polypeptide-oncoprotein 11.66 polypeptide.
                                                                                                                                               WPI; 2002-305565/35
                                                                                                                                                                        мао У,
                                                                                                                                                                                                                      07-JUL-2000; 2000CN-00119427
                                                                                                                                                                                                                                               07-JUL-2000; 2000CN-00119427
                                                                                                                                                                                                                                                                                                CN1333235-A.
                                                                                                                                                                                                                                                                                                                         Homo sapiens
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                                                                                    Example 2; Page 18 (Disclosure); 33pp; Chinese
                                                                                                                                                                                               (SHAN-)
                                                                                                                                                                                                                                                                                                                                                inflammation; PCR; primer.
                                                                                                                                                                                                                                                                                                                                                             haemopathy;
              inflammations.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        179 AGTAGAGATGGAGTTTCTCCATGT 202
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21;
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                                                                                                                                                                        Xie Y;
                                                                                                                                                                                               SHANGHAI BIODOOR GENE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AGTAGAGATGGAGTTTCACATTGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                           (first
            The present DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA;
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              sequence represents
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                                                                                                                        and polynucleotide
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               þ
               human oncogene
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. HIV
                                                   protein
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                            and various
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XC ABL409
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Matches 21
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                                                                                                                                                                                                                  Sequence 24
                                                                                                                                                                                                                                    The invention relates to the human SOX3 protein 13.31 and the polynucleotide encoding it. The polypeptide is used in treating various diseases, such as malignant tumours, heamopathy, HIV infection, immunological diseases and various inflammations. This sequence represents a reverse transcriptase PCR (RT-PCR) primer used in isolation of cDNA encoding the human SOX3 protein 13.31
                                                                                                                                                                                                                                                                                                                                                                                     Mao Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; SOX3 protein 13.31; primer; ss; malignant HIV infection; human immunodeficiency virus; immu inflammation; RT-PCR; reverse transcriptase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human SOX3 protein 13.31 cDNA RT-PCR primer
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                       03-JUL-2002
                                                                                                                                                                                                                                                                                                                    Example
                                                                                                                                                                                                                                                                                                                                         New polypeptide-human SOX3 protein 13.31.
                                                                                                                                                                                                                                                                                                                                                               WPI; 2002-708125/77.
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 Polypeptide-hexokinase protein cDNA isolating primer 2.
                                              ABL40967;
                                                                   ABL40967 standard;
                                                                                                                                                                                                                                                                                                                                                                                                           (BODE-)
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                                                                                                                                                868
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        181 TAGAGATGGAGTTTCTCCATGTTG 204
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                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                   2; Page 16
                                                                                                                                          GGATTACAGGCGTGAGCCACCACG 891
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                                                                                                                          GGATTACAAGCATGAGCCACCATG 24
                                                                                                                                                                                                              BP; 8 A; 6 C; 6 G; 4 T; 0 U; 0 Other;
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                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                      (first
                                                                   DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA;
                                                                                                                                                                                                                                                                                                                 (Disclosure); 32pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1.9%;
                       entry)
                                                                                                                                                                                1.9%;
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Pred. No. 1.4e
0; Mismatches
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Pred. No. 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        .4e+03;
                                                                                                                                                                                .4e+03
                                                                                                                                                                                             DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       nant tumour; haemopathy; immunological disease;
                                                                                                                                                                                         Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 24;
                                                                                                                                                                      Indels
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RESULT 649
ABS57191
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to a novel polypeptide-hexokinase protein. The protein can be expressed by standard recombinant methodology. The novel polypeptide and encoding polypucleotides are used in diagnosis and treatment of malignant tumour, haemopathy, human immunodeficiency virus (HIV) infection, immunological diseases and various inflammations. The present sequence represents the polypeptide-hexokinase protein cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mao
Mao Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                30-JAN-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         diagnosis and treatment of malignant tumors, hemopathy, humaimmunodeficiency virus infection, immunological diseases and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Polypeptide-hexokinase protein and encoding polynucleotide, undiagnosis and treatment of malignant tumors, hemopathy, human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  07-JUL-2000; 2000CN-00117013
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Polypeptide-hexokinase protein; cytostatic; haemostatic; virucide; immunomodulatory; antiinflammatory; RT-PCR; primer; ss.
                                                                                     29-SEP-2000; 2000CN-00125588.
                                                                                                                                 29-SEP-2000;
                                                                                                                                                                             24-APR-2002
                                                                                                                                                                                                                         CN1345971-A
                                                                                                                                                                                                                                                                     Unidentified.
                                                                                                                                                                                                                                                                                                            reverse transcription;
                                                                                                                                                                                                                                                                                                                                    Amylase 9.35; RT-PCR; ss; tumour; haemopathy; antagonist; HIV; human immunodeficiency virus; immunological disease; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                    Amylase 9.35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABS57191 standard;
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                                         (SHAN-) SHANGHAI BIOWINDOW GENE DEV INC
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                                                                                                                                                                                                                                                                                                                                                                                                    specific RT-PCR primer,
                                                                                                                                 2000CN-00125588
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                                                                                                                                                                                                                                                                                                            primer
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Pred. No. 1.4e+03;
0; Mismatches 3
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                                                                                                                                                                                                                                                                                                                                                                                                       #2
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밁 S

WPI;

2002-539369/58

New polypeptide-amylase 9.35 human immunodeficiency virus

for treating malignant tumor, hemopathy, infection, immunological disease and var

and various

inflammations.

Example 2; Page 15

(disclosure); 31pp;

Chinese

coding for the polypeptide and method for producing this polypeptide by using DNA recombination technology. The invention also discloses the method for curing several diseases, such as malignant tumour, haemopathy, human immunodeficiency virus (HIV) infection, immunological diseases and various inflammations by using the polypeptide. The invention also discloses an antagonist for resisting said polypeptide and its therapeutic action and also discloses the application of the polymucleotide for coding this novel amylase 9.35. The sequence presented is the reverse transcription (RT)-PCR primer, #2, which was used to isolate amylase 9.35 cDNA

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RESULT 650
ABQ81233/C
ID ABQ81233/C
ID ABQ812
XX ABQ812
XX ABQ812
XX O5-DEC
XX Human;
KW Human;
KW anorec
KW probe;
XX Homo s
XX Homo;
C6-FEE
XX Homo s
XX H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2002-698629/75.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human 14273 probe.
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The present sequence is a probe, created by PCR, for human 14273 (see ABQ81226), a nucleic acid associated with metabolic disorders. The probe was used to examine the expression profile of human 14273 in different tissues. It was found that 14273 molecules are expressed at high levels.
                                                                                                                                                                                                                                                                                               Identifying a nucleic acid associated with a metabolic disorder, useful for diagnosing metabolic disorders, e.g. obesity, comprises contacting the sample with a probe comprising at least 25 contiguous nucleotides of the 14273 gene.
                                                                                                                                                                                                                 Example 1; Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (MILL-) MILLENNIUM PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              antidiabetic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Έ
                                                                                                                                                                                                                 61; 95pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  disorder; obesity; diabetes; anorexia; cachexia;
; anabolic; transgenic animal; gene therapy;
                                                                                                                                                                                                             English
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of.

syndrome associated protein 11.11, and the polynucleotide encoding it. Also described is a process for preparing the polypeptide by DNA recombination and the application of the polypeptide and polynucleotide in treating various diseases such as cancer and human immunodeficiency virus (HIV) infection. Antagonist against the polypeptide can also be used in treating such diseases. The present sequence for reverse transcriptase (RT)-PCR primer #1 is used with RT-PCR primer #2 (ABK50652 for isolating cDNA encoding human Parkinson's syndrome associated proteins.

(ABK50652)

The present invention relates to the isolation of human

Parkinson's

Example 2; Page 17 (disclosure);

34pp; Chinese

Sequence

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Other;

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RESULT 651
ABK50651/c
IID ABK506
XX ABK506
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Matches 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human, Parkinson's syndrome associated protein 11.11; cancer; human immunodeficiency virus; HIV infection; reverse transcri
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 24 BP; 5 A; 6 C; 8 G; 5 T;
                                                                                                                                                                                                                                                                                                                                                                                                                         New polypeptide-human treating diseases such
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RT-PCR; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human Parkinson's syndrome associated protein 11.11,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      30-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABK50651;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABK50651 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           мао Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       30-JUN-2000; 2000CN-00116961
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (BODE-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2002-292874/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             956 GCAATGGCCAAATCTCGGCTCACT 979
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Xie Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BODE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                         n Parkinsons :
h as cancer a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CO LTD SHANGHAI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          °;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pred.
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                                                                                                                                                                                                                                                                                                                                                                                                                               syndrome associated protein 11.11 and human immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 24;
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924 ATGGAATCTCACTCTGTTACCCAG 947

ABQ81233

standard; DNA;

24 BP

Matches Query Match Best Local

21;

Conservative

0; Mismatches

Pred. Score 19.2;

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.4e+03;

Length 24; Indels

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Similarity

1.9%;

Sequence 24 BP; 4 A; 8 C; 6

G; 6 T; 0

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0 DB 1; Other;

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RESULT 652
ABA04729/c
ID ABA047
XX Human;
XW Human;
XW Human;
XW Antiir
XW Haemog
XX Homo &
XX WO2001
XX WO2001
XX WO2001
XX WO2001
XX WO2001
XX WO2001
XX WANAN
XX WANAN
XX WANAN
PA (SHAN-
XX WAPI;
XX C G AAM477
CC 
RESULT 653
ABQ77823
ID ABQ778
XX
AC ABQ778
AC ABQ778
XX
DT 20-DEC
XX
DT 20-DEC
XX
DE Human
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Best Local Similarity
Matches 21; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 21; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to human ubiquitin-binding enzyme 9 (see AAM47737). The enzyme and its coding sequence are useful in the diagnos and treatment of malignant tumours, haemopathy, HIV infection, immunological diseases and various inflammations. The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human ubiquitin-binding enzyme 9A and encoding polynucleotide, diagnosis and treatment of malignant tumors, hemopathy, human immunodeficiency virus infection, immunological diseases and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; cytostatic; haemostatic; virucide; immunomodulatory; PCR antiinflammatory; gene therapy; ubiquitin-binding enzyme 9; tumchaemopathy; HIV infection; immunological disease; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-MAR-2002
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  Human protein phosphatase 13.64 RT-PCR primer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 24 BP; 3 A; 9 C; 9 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       мао у,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 09-MAY-2000; 2000CN-00115634.
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                                                                                                                                                              ABQ77823 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 2; Page 18; 35pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200188142-A1
                                                       20-DEC-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (SHAN-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PCR Primer, which was used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2002-055699/07.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            925
                                                                                                                                                                                                                                                                                                                                                     874 CAGGCGTGAGCCACCACGCCCGGC 897
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                                                                                                                                                                                                                                                                                                CAGGCGTGAGCCACTGTGCCCGGC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TGGAGTCTCACTCTTTTGCCCAGG 1
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nilarity 87.5%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                    (first entry)
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Pred. No. 1.4e+03;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                Score 19.2;
Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                             .4e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 1;
  SEQ ID NO: 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length
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diagnosis
                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New human protein phosphatase 13.64 polypeptide for treating female genital organ maldevelopment, female sex characteristics abnormality, female genital system tumor and estrogenic relative metabolic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; protein phosphatase 13.64; recombinant production; gene therapy; female genital development disorder; abnormal female sex characteristic; female genital tract tumour; oestrogen-related metabolic abnormality; cytostatic; gynaecological; reverse transcription-PCR; RT-PCR; primer;
                                                                                                                                                                        Human; optineurin; ds; ophthalmological; single nucleotide polymorphism; SNP; glaucoma; progressive ocular hypertensive disorder; glaucoma related disorder; motif; repeat element; regulatory region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 24 BP; 4 A; 3 C; 9 G; 8 T; 0 U; 0 Other;
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                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                         Optineurin promoter motif, repeat element or regulatory region #118.
                                                                                                                                                                                                                                                                                                                            29-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                           ADE14009
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 2; Page 16 (Disclosure); 32pp; Chinese.
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Similarity 87.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BODE GENE DEV CO LTD SHANGHAI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GTAGAGATGGAGTTTCTCCATGTT 203
                                                                                                                                                                                                                                                                                                                                                                                                                        standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GTAGAGATGGGGTTTCACCGTGTT
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                                                                                                                                                                                                                                                                                                                                                                                                                           DNA;
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Pred. No. 1.4e
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 655
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   susceptibility to glaucoma or to a progressive ocular hypertensive disorder resulting in loss of visual field in a patient (or the severity or progression of glaucoma in a patient, comprising providing amplification reaction primers that direct amplification of a selected nucleic acid region containing the variation within the optineurin promoter and amplifying the DNA) and detecting a polymorphism (comprising obtaining a sample containing human genomic DNA, providing a nucleic acid capable of detecting a SNP located within an optineurin promoter, and detecting the polymorphism). The invention is used to diagnose and prognose glaucoma and also to treat glaucoma related disorders. The present sequence is an optineurin promoter motif, repeat element or putative regulatory region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            heterologous sequence, diagnosing or prognosing glaucoma in a sample obtained from a cell or bodily fluid (comprising detecting a polymorphism in a promoter region of the optineurin gene, associated with a glaucoma phenotype), detecting a SNP sequence variation in a sample containing DNA, detecting the presence of an optineurin promoter sequence variation in a sample containing DNA, determining DNA, determining the presence or increased
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to an isolated nucleic acid (N1) comprising at least 20 but not more than 1500 consecutive nucleotides of the optineur promoter appearing as ADE13890. Also included are the optineurin promot operably linked to a heterologous nucleic acid, a nucleic acid capable detecting a single nucleotide polymorphism (SNP) in the optineurin promoter, a host cell comprising the promoter operably linked to a promoter, a host cell comprising the promoter operably linked to a promoter.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             06-MAR-2002; 2002US-00091281.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAT94763 standard; DNA; 19
      US5683885-A
                                                                                                  Synthetic.
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18-FEB-1998
                                                                   Homo sapiens
                                                                                                                                                     antibody; mutation; primer; ss
                                                                                                                                                                                                                                               Human progesterone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1020
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MORISSETTE J.
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                                                                                                                                                                                  progesterone receptor; breast cancer; ovarian cancer; mutant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AGCCTCCCAAGCAGCTGGGATTAC 1043
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                                                                                                                                                                                                                                                                                                           (revised)
(first ent
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                                                                                                                                                                                                                                                  receptor gene primer.
                                                                                                                                                                                                                                                                                                           entry)
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Pred. No. 1
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Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention has developed a novel mechod the discrete invention has developed a novel mechod throlves assaying increased risk for breast or ovarian cancer. The method involves assaying a sample containing human progesterone receptor protein (hpr) with an antibody that distinguishes wild-type hpr from a mutant hPr having a Valco-Leu substitution at amino acid 660, where the presence of such a mutant indicates an increased risk for breast or ovarian cancer. Detection of the G-to-T point mutation gives an odds ratio for ovarian cancer of 3.1 (sensitivity 46%, specificity 78%) and an odds ratio for breast cancer of 2.0 (sensitivity 36%, specificity 78%). (Updated on 25-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     04-NOV-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present sequence represents a primer used in the detection of insertion sequence in intron G in the human progesterone receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Col 7; 26pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Diagnosis of increased risk for breast or ovarian cancer -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                for mutant
                                                                                                                                                             08-JUL-1997.
                                                                                                                                                                                        US5645995-A.
                                                                                                                                                                                                                                                      Breast; ovarian;
detection; point
                                                                                                                                                                                                                                                                                               FISH primer
                                                                                                                                                                                                                                                                                                                                                       AAT84754;
                                                                                                                                                                                                                                                                                                                                                                                AAT84754 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              insertion sequence in intron G in the human progesterone receptor gen
the present invention has developed a novel method for diagnosing an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (BAYU ) BAYLOR
                                                    Kieback
                                                                                                        12-APR-1996;
                                                                                                                                  12-APR-1996;
                                                                              (BAYU ) BAYLOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
nes 19; Conserv
                                                                                                                                                                                                                                            insertion;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       389 AAAGTGCTGGGATTACAGG 407
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               progesterone receptor protein.
                                                                                                                                                                                                                                           arian; cancer; diagnosis; risk; predisposition; human
point mutation; progesterone; receptor; FISH; primen
ion; intron G; fluorescent in situ hybridisation; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             B₽;
                                                                                                                                                                                                                                                                                               for human progesterone receptor intron
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                                                                               COLLEGE MEDICINE
                                                                                                          96US-00629939.
                                                                                                                                   96US-00629939
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Pred. No.
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1.2e+03;
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Diagnosis of increased risk of breast or ovarian cancer - by detecting

WPI; 1997-362926/33

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Increased risk of breast or ovarian cancer can be diagnosed by detecting CC a G to T point mutation at the 1st nucleotide of codon 660 in exon 4 of CC the human progesterome receptor (PR) gene, i.e. nucleotide 2153 of CC AAT84747. The odds ratio is 3.1 (sensitivity 46%, specificity 78%) for CC ovarian cancer, and 2.0 (sensitivity 36%, specificity 78%) for breast CC cancer. The method may also include detecting a C to T point mutation at Ct the 3rd nucleotide of codon 770 in exon 5 of the human PR gene, i.e. CC nucleotide 2485 of AAT84747, and/or an Alu insertion in codon 897 of CC intron G, i.e. AAT84749 inserted between nucleotides 120 and 121 of CC CC concert and with BsrI, and detecting the loss of a BsrI restriction CC site. The mutation in exon 5 can be detected by digesting a test nucleic acid with NIaIII and detecting the loss of a NIaIII restriction site. The mutation in intron G can be detected by digesting a test nucleic acid with NIAIII and detecting the addition of a NIaIII restriction site. The mutation in intron G can be detected by digesting a test nucleic acid with TaqI and detecting the addition of a NIaIII restriction site. The mutation in intron G can be detected by digesting a test nucleic acid with TaqI and detecting the Addition of a NIaIII restriction site. The mutation in intron G can be detected by digesting a test nucleic acid with TaqI and detecting the primers because of the primers are probable of the primers of the primers are probable of the primers of the primers are probable of the pr
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Best Local S
Matches 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200129262-A2
                                                                                         New genotyping oligonucleotide, useful for detecting the presence absence or identity of single polynucleotide polymorphism in a nu
                                                                                                                                                                                                                         Picoult-Newburg L,
                                                                                                                                                                                                                                                                                                                             15-OCT-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                   26-APR-2001
                                                                                                                                                                                                                                                                           (ORCH-) ORCHID BIOSCIENCES INC.
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                                                                      sample.
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                                                                                              polynucleotide polymorphism in a nucleic
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Claim 1; Page 56; 83pp; English

Claim 1; Page 56; 83pp; English

New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nuc

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Single nucleotide polymorphism;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAH38469 standard; DNA; 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15-OCT-1999;
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 inflammation, cancer, nervous system diseases and infection by pathogenic microorganism. The method is also useful in forensic investigations and paternity analysis. The present sequence represents a PCR primer specific for a human can can accept the present sequence represents a PCR primer specific for a human can can be set that the present sequence represents a PCR primer specific for a human can can be set that the present sequence represents a PCR primer specific for a human can can be set that the present sequence represents a PCR primer specific for a human can can be set that the present sequence represents a PCR primer specific for a human can be set to be set to be sequenced as the present sequence represents a PCR primer specific for a human can be set to be set to
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                                                                                                                                                                        Determining wether a human subject has or is at risk of developing (early -onset) Alzheimer's disease comprises detecting the presence/absence of a
                                                                                                                                                                                                                                                                      WPI; 2001-071402/08.
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The present invention describes a method or susceptibility to Alzheimer's disease

for determining the presence of in humans, involving detecting

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The present invention describes a method of arraying genome clones. The compliance (a) clones of the genomic libraries contained in clones in the genomic libraries contained in clones multiwell plates; (b) a primer designed based on the chromosome marker contained is added to the mixture to carry out an amplification reaction; (c) a signal corresponding to the marker is detected from the resultant camplified product to specify the discrimination Nos. of the multiwell confidence is changed so that the same discrimination Nos. succeed to the maximum in the specified discrimination Nos. to array the multiwell confidence; (e) the clones in the multiwell plates of the specified confidence; (e) the clones in the multiwell plates of the specified confidence; (e) the clones in the multiwell plates of the specified confidence; (f) the mixed clones are an array the multiwell confidence; (e) the mixed clones are cultured and the confidence of the specified confidence of the specified plates; (f) the mixed clones are cultured and the confidence of the specified plates are specified from the amplified products; (h) the clones in the multiwell confidence of the specified specified products; (h) the clones in the multiwell confidence are specified from the detected result; and (i) the clones are mixed respected result; and (i) the clones are propertied for multiwell confidence and spain and shadassate of the specified confidence and spain and shadassate of the specified confidence are constituted as the positions on the chromosome and arrayed. The confidence are specified for multiwell confidence are specified to the specified confidence are specified to the specified confidence are specified to the specified specified confidence are specified to the specified specified confidence are specified to the specified specified specified specified specified specified specified specified specified specified
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Alzheimer's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human chromosome 1p36-35 PCR primer SEQ ID NO:1518
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               11-APR-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 4; Page 34; 528pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Arraying genome clones
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2002-144136/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10-MAR-2000; 2000JP-00066716
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12-MAR-2001; 2001JP-00068285
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PCR primer; 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; chromosome
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GENOTEX YG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP; 4 A; 8 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1p36-35; chromosome 21q22.1; genetic analysis; genome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%;
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Pred. No.
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RESULT 661
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                                                                                                                                                  disease, multiple sclerosis, allergy, asthma and diabetic mellitus), diseases or disorders of the immune system, hypersensitivity, anaphylaxis, and blood group incompatibility. The present DNA sequence represents a PCR primer that was used to amplify an intestinal epithelium/peyer's patch M cell-associated DNA sequence of the invention
                                                                                                                                                                                                                                                                               The invention comprises DNA sequences which are associated with intestinal epithelium and peyer's patch M cells. The DNA sequences of the invention are useful for assessing, modifying, modulating or regulating intestinal epithelium or M cell development. The DNA sequences of the
                                                                                                                                                                                                                                                                                                                                                                                          Novel isolated or purified polypeptide encoded by genes associated with intestinal epithelium or M cell development, differentiation or function, useful for treating autoimmune diseases and infectious diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                04-APR-2002; 2002WO-US010873
                                                                                                                        Sequence 19
                                                                                                                                                                                                                              invention are also useful in the treatment of: inflammatory bowel disease, glutenenteropathy, infectious diseases, autoimmune diseases (e.g. haemolytic anaemia, rheumatoid arthritis, dermatitis, Grave's
                                                                                                                                                                                                                                                                                                                                                             Disclosure; SEQ ID NO 543; 152pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-075470/07.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Grave's disease; multiple sclerosis; allergy; asthma; diabetic mellitus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Intestinal epithelium/peyer's patch M cell-associated PCR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    intestinal epithelium cell development; peyer's patch M cell development; inflammatory bowel disease; glutenenteropathy; infectious disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (DIGI-) DIGITAL GENE TECHNOLOGIES INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       autoimmune disease; haemolytic anaemia; rheumatoid arthritis; dermatitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       mmune system disorder; hypersensitivity, lood group incompatibility; ss; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local
                             638
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                                                                          Similarity
                             TGTCACCCAGGCTGGAGTG 656
TGTCACCCAGGCTGGAGTG 19
                                                                                                                      BP; 3 A; 5 C; 7 G; 4 T; 0 U; 0 Other;
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                                                            Conservative
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                                                                         1.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     O'mahony DJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        hypersensitivity; anaphylaxis; lity; ss; human; PCR; primer.
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                                                                         Score 19;
Pred. No.
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                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Evans CF,
                                                                         DB 1; Lo. 1.2e+03;
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                                                                                       Length 19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SP,
                                                                                                                                                     of the invention
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RESULT 662

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RESULT 663
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to a novel method for detecting a gene polymorphism in a human interleukin (II)-18 gene. The method involves detecting a 9 base insertion between -6311 position and -6310 position, a polymorphism at positions -5990, -5316, -4762, -4675, -3268, -689 and -640 of a polymucleotide which consists of a fully defined sequence of 6640 base pairs as given in the specification, where in the 6640bp polynucleotide, the position 6575 is set to +1 from which numbering is performed. The method is useful for detecting gene polymorphism in II-18 gene of human and for detecting adult onset still disease. This polynucleotide sequence represents a probe of the human interleukin-18 gene of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADM32300 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 19 BP; 4 A; 5 C; 7 G; 3 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Detecting gene polymorphism in interleukin-18 gene of human, useful for detecting adult onset still disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           single nucleotide
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                                                                                         Homo sapiens
                                                                                                                                                                               Cytostatic; Gene therapy; breast cancer; human; DLG1; KIAA0783; DPF3; CENPC1; SNP; single nucleotide polymorphism; centromere protein C1;
                                                                                                                                                                                                                                                                        CENPC1 extend primer #59.
                                                                                                                                                                                                                                                                                                                                        26-AUG-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AD080008
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HYUBITTO GENOMICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  standard;
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                                                                                                                                                      P; single nucleotide polymorphism; centromere autoantigen C1; chromosome 4q12-q13.3; extend
                                                                                                                                                                                                                                                                                                                                     (first entry)
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RESULT 6
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AAVS 7
AC AAV
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CT 10-
CT 10-
CX LRP
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KW 1n8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CC risk of breast cancer. The method for identifying a subject at CC risk of breast cancer. The method comprising detecting the presence or CC absence of one or more polymorphic variations associated with breast CC cancer in a nucleic acid sample from a subject. The nucleic acid sample CC cancer in a nucleic acid sample from a subject. The nucleic acid sample CC cancer in a nucleic acid sample (ADO79402), KIAA0783 region (ADO79403), DPF3 CC large homolog 1 (Drosophila)) is also known as synapse-associated protein CC 97, hdlg or SAP97. DLG1 has been mapped to chromosomal position 3q29. The CC gene KIAA0783 is also known as synapse-associated protein is a CC gene kiaA0783 is also known as PHF14 and PHD finger protein 14. KIAA0783 CC has been mapped to chromosomal position 7p21.3. The KIAA0783 protein is a CC it likely to be a transcription factor. The gene DPF3 (D4, zinc and CC double PHD fingers, family 3) is also known as CERP4, cer-44, FL414079 CC and 2810403B03Rik. DPF3 is a Rho family guanine-nucleotide exchange CC gene CENPC1 (centromere protein C1) is also known as Centromere CC (instrumere protein C1) is also known as Centromere CC (instrumere protein C1) is also known as Centromere CC (instrumere protein C1) is also known as Centromere CC (instrumere protein C1) is also known as Centromere CC (instrumere protein C1) is also known as Centromere CC (instrumere protein C1) is also known as Centromere CC (instrumere protein C1) is also known as Centromere CC (instrumere protein C1) is also known as Centromere CC (instrumere protein C1) is also known as Centromere CC (instrumere protein C1) is also known as Centromere CC (instrumere protein C1) is also known as Centromere CC (instrumere protein C1) is also known as Centromere CC (instrumere and a timely transition to anaphase. The method is CC (instrumere protein C1) is also known as Centromere considered to a breast cancer treatment, and in clinical drug trials. The present ion and treatment of breast cancer, to analyze and cCC trials. The present in an exam
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Identifying a subject at risk of breast cancer by detecting the presence of polymorphic variations in the DLG1, KIAA0783, DPF3 or CENPC1 regions which are associated with breast cancer in a nucleic acid sample from a
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24-JUL-2003; 2003US-0490234P.
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                                                                                  LRP5; LDL-receptor related protein; LRP-3; IDDM; diagnosis; endocytosis; insulin dependent diabetes mellitus; autoimmune disease; glomerulonephritis; inflammation; viral infection; osteoporosis;
                                                                                                                                                                                                                  10-FEB-1999
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                                          PCR primer;
                                                                 ypercholesterolemia; Alzheimer's disease;
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                                                                     low density lipoprotein;
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention describes LRP5 (low density lipoprotein (LDL) receptor related protein, previously designated LRP-3). AAV85587 to AAV85822 represent exon primers used for obtaining LRP5 CDNA. Nucleic acid molecules (NAMS) encoding LRP5 can be used for determining if an individual is succeptible to insulin dependent diabetes mellitus (IDDM). The NAMS or proteins can be used for reducing triglyceride levels in the serum of an individual. Therapies that affect LRP5 may also be useful in
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05-JUN-1997;
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                                                                         hypercholesterolemia; Alzheimer's PCR primer; ss.
                                                                                                LRP5; LDL-receptor related protein; LRP-3; IDDM; diagnosis; endocytosis; insulin dependent diabetes mellitus; autoimmune disease; glomerulonephritis; inflammation; viral infection; osteoporosis;
                                                                                                                                                  LRP5 SNP primer 58-7
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IR, Metzker ML, Nakagawa
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                                                                                                                                                                                                                                                                                                                     Human mdm2 gene; proliferation; tumour; phosphorothicate; p53; cancer; antisense; modulation; oligonucleotide; expression; inhibition; hyperproliferation; blood cancer; brain cancer; breast cancer; lung cancer; soft tissue cancer; psoriasis; fibrosis; atherosclerosis;
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Hey P, Kawaguchi Y, Mer
Phillips MS, Twells RCJ;
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05-JUN-1997;
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100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1.2e+03;
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kagawa Y;
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Matches 19
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                                                                                                                                                                                                                                                                                                                                                                            07-JAN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                            AAZ37720 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 3 A; 10 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New antisense compounds used to treat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Miraglia LJ,
                                                                                                                                      26-MAR-1998;
                                                                                                                                                                26-MAR-1999;
                                                                                                                                                                                        30-SEP-1999
                                                                                                                                                                                                                                                    Synthetic
                                                                                                                                                                                                                                                                            restenosis; ss.
                                                                                                                                                                                                                                                                                       hyperproliferation; blood cancer lung cancer; soft tissue cancer;
                                                                                                                                                                                                                                                                                                                          Human mdm2 gene;
                                                                                                                                                                                                                                                                                                                                                   Human mdm2
                                                                                                                                                                                                                                                                                                                                                                                                     AAZ37720;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1999-610754/52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       26-MAR-1998;
                                                              WPI; 1999-610754/52
                                                                                                                                                                                                              WO9949065-A1
                                                                                                                                                                                                                                                                                                               antisense;
                                                                                                               (ISIS-) ISIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
                                                                                                                                                                                                                                       sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 644 CCAGGCTGGAGTGCAGTGG 662
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CCAGGCTGGAGTGCAGTGG 2
                                                                                                                                                                                                                                                                                                    gene; proliferation; tumour; phosphorothioate; p53;
modulation; oligonucleotide; expression; inhibition;
feration; blood cancer; brain cancer; breast cancer;
                                                                                                                                                                                                                                                                                                                                                  phosphorothicate oligodeoxynucleotide #250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Page
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                                                                                                               PHARM
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                                                                                     Nero P,
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                                                                                                                                      98US-00048810.
                                                                                                                                                                99WO-US006702
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                                                                                                               INC
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                                                                                      Graham MJ,
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                                                                                      Monia
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      hyperproliferative conditions.
                                                                                                                                                                                                                                                                                        fibrosis; atherosclerosis;
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Example 9; Page 54; 157pp; English

New antisense compounds used to treat

eg.

hyperproliferative conditions.

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RESULT 668
AAF80874/c
ID AAF808
XX AAF808
XX AAF808
XX AAF808
XX AAF808
XX Human
XX Antiss
OS Homo s
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PD 06-FBE
XX 26-MAF
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           hyperproliferative gene expression. The antisense compound is used to treat an animal having a disease or condition associated with mdm2, particularly a hyperproliferative condition, more particularly cancer, especially of the blood, brain, breast, lung or soft tissue, or psoriasis, fibrosis, atherosclerosis or restenosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel antisense compound 8-30 nucleobases in leacid molecule encoding human mdm-2 useful for mof human mdm-2 and reducing hyperproliferation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human mdm2 phosphorothioate oligonucleotide
                                                                                                                                                                                 The present invention relates to an antisense compound 8-30 nucleobases in length targeted to nucleobases 1-308 of the 5' untranslated region, 1776-1806 of the translation termination codon region or 1818-2370 of the 3' untranslated region of a nucleic acid molecule encoding human mdm-2. The invention is useful for reducing hyperproliferation of human cells, modulating the expression of mdm2 in human cells or tissues or in vitro.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              02-MAY-2001
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                                                                                                                                                                                                                                                                                                                                                                                      Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-190948/19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Miraglia LJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               26-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   26-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Antisense; mdm2;
                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (ISIS-) ISIS
                                                                                                                                                               hyperproliferative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    536 TCCTGCCTCAGCCTCCCAA 554
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                                                                                                                                                                                                                                                                                                                                                                                      9; Col 33; 77pp; English.
                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TCCTGCCTCAGCCTCCCAA 2
                                                                                                           20 BP; 5 A; 2 C; 10 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP; 5 A; 2 C; 10 G; 3 T; 0 U; 0 Other;
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Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nero P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               98US-00048810
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                             100.0%;
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                                                                                                                                                                  disorder includes cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Graham MJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20
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Pred. No.
                             Score 19;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
  Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1;
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                                                       DB 1;
                               1.2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                              in length targeted to a nucleic for modulating the expression ation of human cells.
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                                                          Length 20;
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RESULT 669
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                                                                                                                          The present invention relates to an antisense compound 8-30 nucleobases in length targeted to nucleobases 1-308 of the 5' untranslated region, 1776-1806 of the translation termination codon region or 1818-2370 of the 3' untranslated region of a nucleoic acid molecule encoding human mdm-2. The invention is useful for reducing hyperproliferation of human cells, modulating the expression of mdm2 in human cells or tissues or in vitro.
                                                                                                                                                                                                                                 Novel antisense, compound 8-30 nucleobases in length targeted to a nucleoacid molecule encoding human mdm-2 useful for modulating the expression of human mdm-2 and reducing hyperproliferation of human cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAF80867 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human mdm2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     02-MAY-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAF80867;
                                                                                          Sequence
                                                                                                                                                                                                          Example 9; Col 31; 77pp; English.
                                                                                                                                                                                                                                                                                                        Miraglia LJ,
                                                                                                                                                                                                                                                                                                                                                      26-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                            26-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Antisense; mdm2; hyperproliferation; cancer; psoriasis;
                                                                                                                                                                                                                                                                                  WPI; 2001-190948/19
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                 The hyperproliferative disorder includes cancer or
                                                                                                                                                                                                                                                                                                                               (ISIS-)
                     644 CCAGGCTGGAGTGCAGTGG
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                                              19;
                                                                                                                                                                                                                                                                                                                                ISIS
                                                         Similarity
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                                                                                          20
CCAGGCTGGAGTGCAGTGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              phosphorothicate oligonucleotide
                                                                                          BP; 3 A;
                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                PHARM INC
                                                                                                                                                                                                                                                                                                        Nero P,
                                                                                                                                                                                                                                                                                                                                                        98US-00048810
                                                                                                                                                                                                                                                                                                                                                                              99US-00280805
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA;
                                                         1.9%;
                                                                                          10 C; 4 G; 3 T; 0 U; 0 Other;
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                                                                    Score 19;
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                                                           Pred.
                                               Mismatches
                                                                                                                                                                                                                                                                                                          Monia BP,
                                                           No.
                                                                     DB 1;
                                                          1.2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 #241.
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                                                                     Length 20;
                                              Indels
                                                                                                                   psoriasis
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536 TCCTGCCTCAGCCTCCCAA 554
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EP1054059-A1

Catenin-binding

zinc finger protein; cancer; neurological disorder;

PCR

primer FVR510F.

screening;

PCR primer;

88

Human catenin-binding zinc finger protein

XXXX

07-MAR-2001 AAC88720;

(first

entry)

AAC88720 RESULT 670

AAC88720 standard;

DNA;

20

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ARBSULT 671
AAD12635
ID AAD1266
XX AAD126
XX AAD126
XX AAD126
XX Human
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XX Human
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention is related to the coding sequence and protein fragments of a human catenin-binding zinc finger protein. The coding sequence was isolated from a human kidney cDNA library, but is expressed in most human tissue. The sequences provided by the invention can be used in the diagnosis and treatment of cancer and neurological disorders, and in drug screening to identify compounds capable of the same
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Van
Disclosure; Page 66; 160pp; English.
                                              Novel recombinant nucleic acids useful for diagnosing, prognosing and/or treating cancer and neurological disorders, corresponds to a protein binding to alpha-catenin protein and with signal transduction function.
                                                                                                                                                                                                                                                                                                                                                                 18-MAY-2000; 2000WO-EP004535
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; ANC_2H01 protein; catenin-binding protein; signal transduction; gene regulation; zinc finger protein; alphaN-catenin; drug screening; therapy; cancer; neurological disorder; cytostatic; neuroprotective;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human ANC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAD12635;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nucleic acid or its fragments, useful for diagnosing and treating cancer and neurological disorders, corresponds to a catenin-binding protein in signal transduction and gene regulatory pathways.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 5 A; 3 C; 8 G; 4 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Page 17; 71pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                       99EP-00204512.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CDNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       sequencing forward primer, FVR510F.
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Pred. No.
                                                                                                                                                                                                                 Janssens
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Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to human catenin-binding proteins and their corresponding cDNA molecules which functions in signal transduction and gene regulatory pathways. The invention also provides an isolated and/or recombinant nucleic acid or its functional fragment, homologue or derivative, corresponding to a alpha-catenin binding protein. The invention also relates to a novel human zinc finger protein binding with a member of the a-catulin/vinculin family, preferably with a human isoform of alpha N-catenin (neural form). The invention also relates to
                                                                                                                                                                                                                                                                                                                                                                                                                                Human; mdm2; hyperproliferative disorder; cancer; psoriasis;
atherosclerosis; tumour; cytostatic; anti psoriatic;
anti arteriosclerotic; vasotropic; antisense; phosphorothioate;
                                                                                                                                                                                                      26-MAR-1998;
26-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human mdm2 antisense oligonucleotide 31620
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21-NOV-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAS29482 standard; DNA; 20 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               the field of drug discovery, diagnosis, prognosis and treatment of can and neurological disorders. The present sequence is a primer which is
                                                                                 WPI; 2001-535565/59.
                                                                                                         Miraglia
                                                                                                                                          (NERO/)
(GRAH/)
(MONI/)
                                                                                                                                                                                                                                         02-JAN-2001;
                                                                                                                                                                                                                                                                23-AUG-2001.
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                                                                                                                                                                                                                                                                                                                                                                     modified_base
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19; Conser
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                                                                                                                              NERO P.
GRAHAM M J.
MONIA B P.
COWSERT L M.
                                                                                                                                                                              MIRAGLIA L J
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20 BP; 5
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                                                                                                                                                                                                                                          2001US-00752983
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                                                                                                         Nero
                                                                                                                                                                                                     98US-00048810.
99US-00280805.
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/mod= "OTHER= All phosphorothioate linkages,
/note= "OTHER= All phosphorothioate linkages,
additionally bases 1-6 and bases 15-20 are 2'.-O-
methoxyethyl bases, and bases 7-14 are deoxynucleotides"
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                                                                                                          Monia BP,
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                                                                                                          Cowsert
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of a nucleic acid encoding

human

mdm2.

reating e.g. translation

cancer, comprises termination codon region)

Example 9;

Page 18; 81pp; English

nucleobases targeted a region (e.g. transla

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RESULT 673
AAS29489/c
ID AAS294
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Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention relates to antisense compounds, 8-30 nucleobases in length targeted to the 5' untranslated region, translation termination codon region, 3' untranslated region, coding region or translation start site of a nucleic acid encoding human mdm2, where the antisense compound modulates the expression of human mdm2. The antisense oligonucleotides of the invention are useful for encoding human mdm2 and for inhibiting the expression of human mdm2. They may be used for treating an animal having a disease or condition associated with amplification of mdm2 gene or overexpression of mdm2 e.g. a hyperproliferative disorder such as cancer (blood, brain, breast, lung, or a soft tissue cancer) and poriasis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        fibrosis, atherosclerosis or restenosis, tumours, colorectal carcinoma and chronic myelogenous leukemia. The antisense compound may be administered with a chemotherapeutic agent to overcome drug resistance. The antisense compound reduces hyperproliferation of human cells. The method, which involves the use of the antisense compound, is also useful for detecting the role of mdmz expression in various cell functions and physiological processes and useful in both clinical research and diagnostic tools. AAS29242-AAS29507 represent the human mdm2 antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; mdm2; hyperproliferative disorder; cancer; psoriasis; atherosclerosis; tumour; cytostatic; anti psoriatic; anti arteriosclerotic; vasotropic; antisense; phosphorothioa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20 BP; 3 A; 10 C; 4 G; 3 T; 0 U; 0 Other;
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                                                                                                      (MIRA/)
(NERO/)
(GRAH/)
                                                                                                                                                                            26-MAR-1998;
26-MAR-1999;
                                                                                                                                                                                                                                02-JAN-2001; 2001US-00752983
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                                                                                                                                                                                                                                                                                                                                                                                                                               modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human mdm2 antisense oligonucleotide 31784.
 WPI; 2001-535565/59
                               Miraglia LJ,
                                                                     (MONI/)
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                                                                     MONIA B P.
COWSERT L M.
                                                                                                      NERO P.
GRAHAM M J.
                                                                                                                                      MIRAGLIA L J.
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                                 Nero P,
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99US-00280805
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additionally bases 1-6 and bases 15-20 are 2'-0-
methoxyethyl bases, and bases 7-14 are deoxynucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                             mod_base= OTHER
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                                   Graham MJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20
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Pred. No.
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                                     Cowsert
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An antisense compound, useful for treating e.g. cancer, comprises nucleobases targeted a region (e.g. translation termination codon of a nucleic acid encoding human mdm2. region)

Page 18; 81pp; English.

cc site of a nucleic acid encoding human mdm2, where the antisense compound cc modulates the expression of human mdm2. The antisense of incompound condulates the expression of human mdm2. The antisense oligonucleotides of the invention are useful for encoding human mdm2 and for inhibiting the expression of human mdm2. They may be used for treating an animal having care disease or condition associated with amplification animal having coverexpression of mdm2 e.g. a hyperproliferative disorder such as cancer (blood, brain, breast, lung, or a soft tissue cancer) and psoriasis, atherosclerosis or restenosis, tumours, colorectal carcinoma cardinomic myelogenous leukemia. The antisense compound may be cadministered with a chemotherapeutic agent to overcome drug resistance. The antisense compound, is also useful comethod, which involves the use of the antisense compound, is also useful conditions and compound the role of mdm2 expression in various cell functions and conditions are conditions and conditions and conditions are conditions. AAS29242-AAS29507 represent the human mdm2 antisense conditions are conditions. The present invention relates to antisense compounds, length targeted to the 5' untranslated region, translated codon region, 3' untranslated region, coding region or site of a nucleic acid encoding human mdm2, where the oligonucleotides of the present invention translation termination or translation start 8-30 nucleobases

Sequence 20 BP; 5 A; 2 C; 10 G; 3 T; 0 U; 0 Other;

Ś Query Match Best Local & Matches 536 TCCTGCCTCAGCCTCCCAA 554 19; Similarity Conservative 100.0%; 1.9%; 0; Score 19; Pred. No. Mismatches DB 1; Le 1.2e+03; Length 20 Indels 0 Gaps 0

AAD12408 standard; DNA; 20 ВP 吊

20

TCCTGCCTCAGCCTCCCAA 2

25-SEP-2001 (first entry)

AAD12408;

Human caspase 8 mRNA antisense compound ISIS 107686. Caspase 8; infection; inflammation; flammation; tumour; research reagent; cytostatic; human; phosphorothioate; ss.

gene therapy; Synthetic. Homo sapiens antisense;

RESULT 674
AAD12408/c
ID AAD124
XX AAD124
XX AAD124
XX Caspas
KW Gene ti
XX Gene ti
XX Homo s
OS Synthe
FT modifi
FT modifi modified_base modified_base modified_base modified_base modified_base modified_base /mod_base= m5c Location/Qualifiers /*tag= /note= "Phosphorothioate backbone" /mod_base≃ /mod_base= note= mod mod. *tag= *tag= *tag= base= OTHER e= "2'-methoxyethyl base= OTHER m5c OTHER (2'-MOE) nucleotides"

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RESULT 675
AAL61524
ID AAL615
XX
AC AAL615
XX
AC AAL615
XX
DT 22-SEF
XX
DE Human
XX
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                                                                                                               Query Match
Best Local S
Matches 19
      Human inhibitor-kappa B-R antisense oligonucleotide,
                                                                                                                                            Sequence
                                                                                                                                                        prophylaxis e.g. to prevent or delay infection, inflammation formation, and as a research reagent. The present sequence is antisense compound targetted to human caspase 8 mRNA
                                                                                                                                                                              of human caspase 8. The antisense compound is useful for diagnosing a treating diseases associated with the expression of caspase 8 and for
                                                                                                                                                                                                          Example 15; Col 45-46; 56pp; English.
                                                                                                                                                                                                                      New antisense compounds capable of modulating expression of caspase the diagnoses, prophylaxis and treatment of diseases associated with expression of caspase 8, e.g. inflammation and tumor formation.
                                                                                                                                                                                                                                                                                                            19-JAN-2000; 2000US-00487445.
                                                                                                                                                                                                                                                                                                                                                                   modified_base
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                      22-SEP-2003
                                                 AAL61524 standard;
                                                                                                                                                                                           The invention relates to antisense compounds which inhibit the expression
                                                                                                                                                                                                                                                                                              19-JAN-2000;
                                                                                                                                                                                                                                                                                                                                        US6258600-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  modified_base
                                                                                                                                                                                                                                                                                                                           10-JUL-2001.
                                                                                                                                                                                                                                                     2001-432165/46.
                                                                                                  646
                                                                                     20
                                                                                                                19;
                                                                                                                                                                                                                                                                                 ISIS PHARM INC
                                                                                                                       Similarity
                                                                                                                                            20
                                                                                                  AGGCTGGAGTGCAGTGGCG 664
                                                                                     AGGCTGGAGTGCAGTGGCG 2
                                                                                                                                                                                                                                                                  Cowsert LM;
                                                                                                                                           BP; 4 A; 10 C; 3 G; 3 T; 0 U; 0 Other;
                                                                                                                Conservative
                                                                                                                                                                                                                                                                                              2000US-00487445
                    (first entry)
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nd_base=
"2'
                                                                                                                                                                                                                                                                                                                                                                                                                    /mod_base=
16. .20
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/mod_base=
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                                                                                                                                                                                                                                                                                                                                                                                                                                  *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          mod_base=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 *tag=
                                                 DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     base=
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                                                                                                                       100.0%;
                                                                                                                              1.9%;
                                                                                                                                                                                                                                                                                                                                                                                               "2'-methoxyethyl (2'-MOE) nucleotides"
                                                 20
                                                                                                                                                                                                                                                                                                                                                                           m5c
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                                                 ₽P.
                                                                                                                                                                                                                                                                                                                                                                                                       OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                            mSc
                                                                                                               0
                                                                                                                       Score 19;
Pred. No.
                                                                                                                      Pred.
                                                                                                               Mismatches
                                                                                                                      1.2e+03;
                                                                                                                              DB 1;
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                                                                                                                             Length 20;
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        ISIS #130449
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                                                                                                                                                                        or
                                                                                                               Gaps
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RESULT 676
ADC65799
ID ADC657
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The invention relates to antisense compounds targetted to a nucleic acid molecule encoding human inhibitor-kappa B-R (also known as I-kappaBR, I'KBR, I-kappa-B-related, ikappabr, miclear factor of kappa light polypeptides gene enhancer in B-cells inhibitor-like 2 and NFKBIL2) to inhibit its expression. Antisense compounds of the invention are useful for treating diseases or conditions associated with the expression of increased cytokine expression, or a result of infection (e.g. bacterial, viral or parasitic). They are useful for diagnostics, therapeutics, prophylaxis e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and kits and in distinguishing between functions of various members of a biological pathway. They are also useful in antisense therapy. The present sequence is an oligonuclectide targetted to human inhibitor-kappa B-R DNA
                                                                                               Matches
                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New antisense oligonucleotides targeted to nucleic acids encoding inhibitor-kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a heighte immune response or infection.
                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 3; Page 74; 108pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2003-468635/44.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        13-NOV-2001; 2001US-00993731
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ISIS-) ISIS PHARM
                                                                                                                       Local
                                            645 CAGGCTGGAGTGCAGTGGC 663
μ.
                                                                                             l Similarity
19; Conserv
                                                                                                                                                                                               20 BP; 4
CAGGCTGGAGTGCAGTGGC 19
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                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2002WO-US035597.
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16. .20
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/mod_base=
/note= "2'-
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                                                                                                                                                                                               A; 4 C; 9 G; 3 T; 0 U; 0 Other;
                                                                                                                       1.9%; Score 19;
100.0%; Pred. No
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= "2'-methoxyethyl (2'-MOE)
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                                                                                             0;
                                                                                               Mismatches
                                                                                                                         No.
                                                                                                                       1.2e+03;
                                                                                                                                                 DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (2'-MOE)
                                                                                                                                            Length 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          nucleotides"
                                                                                               Indels
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ADC65799 standard; DNA;

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RESULT 677
ADD21685/c
ID ADD216
XX A
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     18-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADC65799;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention comprises antisense oligonucleotides that are targeted to the nucleic acid encoding transforming growth factor beta (TGF-beta) receptor II. The antisense oligonucleotides of the invention are useful for treating: hyperproliferative disorders (e.g. breast cancer), or an autoimmune disorder (e.g. rheumatoid arthritis). The present DNA sequence represents a 2'-O-methoxyethyl gapmer oligonucleotide with a phosphorothioate backbone that is targeted to human TGF-beta receptor II.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New compound having a sequence targeted to a nucleic acid encoding Transforming growth factor beta-receptor II, useful for preparing a composition for treating hyperproliferative disorder e.g., lung, liver,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      human; antisense oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 03-JAN-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 15; SEQ ID NO 95; 141pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       colon
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19-JUN-2002; 2002WO-US019665
                                                                                                                                                                   antisense oligonucleotide; human; mdm2; hyperproliferation;
hyperproliferative disorder; cancer; psoriasis; fibrosis;
atherosclerosis; restenosis; apoptosis modulation; p21; ss;
                                                                                                                                                                                                                                                                             Human mdm2 antisense oligonucleotide #248.
                                                                                                                                                                                                                                                                                                                                 15-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                              ADD21685
                                                                                   Homo sapiens
                                                                                                                                         2'-methoxyethoxy-residue; phosphorothioate
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     541 CCTCAGCCTCCCAAGTAGC 559
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       or gastric cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    N
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20 BP; 5 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Wyatt JR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PHARM INC
                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        9 C; 3 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            targeted antisense oligonucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 19;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1.2e+03;
hes 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        autoimmune
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
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RESULT 678
ADD21678/c
ID ADD216
XX IONA
PN WO2003
XX IONA
PN IONA
XX INTAG1
PN MITAG1
PN MITAG1
PN WPI; 2
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Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention comprises antisense oligonucleotides which are targeted to the human mdm2 gene. The antisense oligonucleotides of the invention are useful for reducing hyperproliferation of human cells. The antisense oligonucleotides are also useful for treating: hyperproliferative disorders (e.g. cancer), psoriasis, fibrosis, atherosclerosis, or restenosis. The antisense oligonucleotides are also useful for modulating apoptosis, and for increasing expression of p21. The present DNA sequence represents a human mdm2 gene antisense oligonucleotide of the invention. The present sequence contains 2'-methoxyethoxy-residues and has a phosphorothioate backbone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel antisense compound targeted to 5' untranslated region, region, or intron:exon junction of nucleic acid molecule encouseful for treating e.g. cancer, psoriasis or restenosis by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Miraglia LJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        04-DEC-2001; 2001US-00005344.
                                                                                                                                                                                                                                                antisense oligonucleotide; human; mdm2; hyperproliferation; hyperproliferative disorder; cancer; psoriasis; fibrosis; atherosclerosis; restenosis; apoptosis modulation; p21; ss;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 9; SEQ ID NO 250; 289pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  02-DEC-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               12-JUN-2003
                                                                                                                                                                                                                                                                                                       Human mdm2
                                                                                                                                                                                                                                                                                                                                                                                          ADD21678
WPI; 2003-577263/54
                           Miraglia LJ,
Manoharan M;
                                                                                                                         02-DEC-2002;
                                                                                                                                                   12-JUN-2003.
                                                                                                                                                                             WO2003048315-A2
                                                                                                                                                                                                                                    2'-methoxyethoxy-residue;
                                                                                                                                                                                                                                                                                                                                    15-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                ADD21678;
                                                                    (ISIS-) ISIS
                                                                                               04-DEC-2001; 2001US-00005344
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 536 TCCTGCCTCAGCCTCCAA 554
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19; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                        standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20 BP; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TCCTGCCTCAGCCTCCCAA 2
                                                                                                                                                                                                                                                                                                         antisense oligonucleotide #241.
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                                                                     PHARM INC
                                                                                                                           2002WO-US038281.
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                                        Nero
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                                          PS,
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Pred. No.
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1.2e+03;
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e encoding mdm2,
s by inhibiting
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Novel antisense compound targeted to 5' untranslated region, coding region, or intron:exon junction of nucleic acid molecule encoding mdm2, useful for treating e.g. cancer, psoriasis or restenosis by inhibiting

expression.

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ADD25037/c
ID ADD250
XX
ADD25037/c
ID ADD250
AC ADD250
AC ADD250
XX
Caspas
KW Caspas
KW antise
KW naemat
KW naemat
KW naemat
KW cancer
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TH Key
FT modifi
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   antisense gene therapy; apoptosis; hyperproliferative disorder; haematopoletic disorder; autoimmune disorder; viral infection; AIDS; neurological disorder; Alzheimer's disease; Parkinson's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 3 A; 10 C; 4 G; 3 T; 0 U; 0 Other;
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                 19-JAN-2000;
11-JAN-2001;
                                                                                                                                                                                                                                                                                            modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cancer; human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human caspase-8 antisense oligonucleotide ISIS 107686.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15-JAN-2004
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                                                                                                                                                                                                                          modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     amyotrophic lateral sclerosis; retinitis pigmentosa; blood cell disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Caspase-8; cytostatic; immunosuppressant; anti-HIV; ss;
                                                                    12-JUL-2002; 2002US-00181177
                                                                                                                                        US2003083296-A1
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                 2000US-00487445
2001WO-US000955
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/mod_base=
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/note= "Phosphorothicate backbone
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== "2'-methoxyethyl residues"
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                                                                                                                                                                                                                                              residues"
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20

Matches Query Match Best Local

646 19;

AGGCTGGAGTGCAGTGGCG AGGCTGGAGTGCAGTGGCG 2

664

Local Similarity

Conservative

0,

Mismatches

0

Gaps

0

1.9%;

Score 19; Pred. No.

DB 1; 1.2e+03; 0 Length Indels

RESULT 680
ABZ979
ID ABZ979
XX ABZ97
XX ABZ

24-APR-2001; 2001US-0286137P

23-APR-2002; 2002WO-US013135

31-OCT-2002

Homo sapiens

inflammation;

respiratory

disease; ds.

Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;

Human RANTES oligonucleotide sequence.

17-OCT-2003

(first entry)

ABZ97910 standard; DNA; 20

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with at least an 8-nucleobase portion of an active site on a nucleic acid complexue encoding caspase 8, a composition comprising the compound and a composition of caspase 8 in cells or tissues with the compound so that compression of caspase 8 in cells or tissues with the compound so that capression of caspase 8 is inhibited) and treating an animal having a condition associated with caspase 8 by administering to the canimal a therapeutic or prophylactic amount of the compound so that capression of caspase 8 is inhibited. The compound, composition and compended are useful for treating a disease or condition associated with caspase 8, such as hyperproliferative, haematopoietic or autoimmune condition such as hyperproliferative, haematopoietic or autoimmune condition such as hyperproliferative, haematopoietic or autoimmune condition infection such as AIDS, neurological disorders (e.g. Alzheimer's disease, Parkinson's disease, amyotrophic lateral sclerosis, cretinitis pigmentosa), blood cell disorders and cancer. They are also cuseful in research and diagnostics for modulating the expression of conference is a caspase-8 targeting antisense conjugonucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to a compound 8-30 nucleobases in length targeted to, and which specifically hybridises with a nucleic acid molecule encoding caspase 8 (a protein involved in apoptosis) and inhibits the expression of caspase 8, i.e. an antisense oligonucleotide. Also included expression of mucleobases in length that specifically hybridises are a compound 8-30 nucleobases in length that specifically hybridises
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New compounds, particularly antisense oligonucleotides targeted to a nucleic acid encoding caspase 8, useful for treating a disease/condition associated with caspase 8, such as hyperproliferative or autoimmune
     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 15; SEQ ID NO 94; 59pp;
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COWSERT L M.
20 BP; 4 A; 10 C; 3 G; 3 T; 0 U; 0 Other;
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RESULT 681
ABZ98002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cc initiation codon, coding region, 5. or 3. and genomic flanking regions, cc 5. and 3. intron-exon junctions, or regions within 2-10 nucleotides of cc junctions of genes encoding a polypeptide associated with lung and/or cc nasal airway dysfunction and a second active agent comprising an cc antiinflammatory steroid and ubiquinone. A composition of the invention cc nas antiinflammatory, antiallergic, antiasthmatic, hypotensive, cc immunosuppressive, and cytostatic activity. The composition may have a cc use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also cc for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels cf, or reducing sensitivity to adenosine, reducing levels of adenosine creceptor, producing bronchodilation, increasing levels of this patent is net represented in the printed specification, but was obtained in electronic format directly from WIPO at fre, wipo, int/nub/ished nor remembers.
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Matches 19
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Miller S,
                                                                                                                                                                         Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-229219/22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to a novel pharmaceutical composition, first active agent comprising an oligonucleotide antisense to initiation codon, coding region, 5' or 3' end genomic flanking.
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                                                                                                                                                                                                                                                                       Human RANTES oligonucleotide sequence.
                                                                                                                                                                                                                                                                                                       17-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                     ABZ98002 standard;
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                                                               31-OCT-2002
                                                                                                                            Homo sapiens.
                               23-APR-2002; 2002WO-US013135
                                                                                               WO200285308-A2
                                                                                                                                                             inflammation;
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Tang L,
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Shahabuddin
                                                                                                                                                                                                                                                                                                                                                                      DNA;
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to a novel pharmaceutical composition, which has first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, or regions within 2-10 nucleotides of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                          lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; SEQ ID NO 13244; 872pp; English.
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                                                                                                                                                                                                                                                     Sequence 20
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                                                          638 TGTCACCCAGGCTGGAGTG 656
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N
                                                                                                                                                             Similarity
   TGTCACCCAGGCTGGAGTG
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Tang
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L, Shahabuddin
                                                                                                                                                             100.0%;
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                                                                                                                                                             Score 19;
Pred. No.
      20
                                                                                                                               Mismatches
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                                                                                                                                                                1.2e+03;
                                                                                                                                                                                             DB 1;
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                                                                                                                                                                                             Length 20;
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RESULT 682
ABD31033
                                                                                                                                                                                               Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory disease; pulmonary hyportension; respiratory distress syndrome; allergic rhinitis; pulmonary hyportension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis;
                                                                                                                                                                      beta-adrenergic agonist; respiratory disease; pulr respiratory distress syndrome; allergic rhintits; respiratory distress syndrome; allergic rhintits; emphysema; chronic obstructive pulmonary disease; pulmonary transplantation rejection; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                Human RANTES-derived oligonucleotide SEQ ID 13244.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABD31033;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABD31033 standard; DNA; 20
                                                                                                                             sapiens
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24-APR-2001; 2001US-0286137P

23-APR-2002; 2002WO-US013143.

31-OCT-2002. WO200285309-A2

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ABD30941
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CC oligonucleotides are derived from a gene encoding or regulating construction of a target polypeptide associated with lung airway or lung construction or cancer and can be anti-sense to the corresponding mRNA. CC The invention also describes a kit, that comprises: (a) a delivery constructions for adding a carrier and for use of the kit. The composition of instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, confidency of the invention for a manufacture and cytostatic activity, is a composition comprises oligo and is administered to reduce the production composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The conflammation, allergies and/or bronchoconstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vascoconstriction, respiratory conflammation, allergies, asthma, impeded respiration, respiratory conflammation, allergies, asthma, impeded respiration, respiratory construction repection, pulmonary infections, bronchities or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the manufaction in the target RNA serves to prevent the breakdown of the manufaction in the target RNA serves to prevent the breakdown of the manufaction in the target RNA serves to prevent the breakdown of the manufaction and the product of the anti-sense oligos corresponding to the manufaction and the production and productions between the breakdown of the manufaction and the production of the anti-sense oligon to the action the production of the anti-sense oligon to the serves to the serves to the production of the anti-sense oligon to the action of the anti-sense oligon to the production of the anti-sense oligon
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Miller S,
                           Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          bronchoconstriction, respiratory tract reducing adenosine sensitivity, levels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24-APR-2001; 2001US-0286036P
                                                                                                                                                                   Human RANTES-derived oligonucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (EPIG-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            oligonucleotides into products that free adenosine into , lung, brain, heart, kidney, etc, tissue environment an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           638
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Tang
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                unwanted effects due to it
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L, Shahabuddin
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   immunosuppressive; cytostatic; cystic
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Pred. No.
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                                                                                                                                                                   SEQ ID
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1.2e+03;
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                                                                                                                                                                            CC expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. CC The invention also describes a kit, that comprises: (a) a delivery construction or cancer and can be anti-sense to the corresponding mRNA. CC The invention also describes a kit, that comprises: (a) a delivery constructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, consists of the invention has antiallergic, antiinflammatory, antiasthmatic, and beta-adrenergic agonist. The composition is useful for preventing or construction are respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The composition control of target polypeptide present in the lungs. The confilammation, allergies and/or bronchoconstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are associated confilammation, allergies and/or surfactant hypoproduction are associated confilammation, allergies and/or surfactant hypoproduction are associated confilammation, allergies, asthma, impeded respiration, respiratory hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary content of the anti-sense oligos corresponding to the oligonucleotides into products that free adenosine into the system of the production of the content of the content of the system of the content of the content of the system of the content of the content of the system of the content 
                                                                                                          Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptor surfactant depletion or hyposecretion, when administered to a mammal oligonucleotides are derived from a gene encoding or regulating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory diserses syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-093058/08
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                                                                                                                                                                                                                    the oligonucleotides into products that e.g., lung, brain, heart, kidney, etc, to prevent any unwanted effects due to it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15; SEQ ID NO 13152;
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N
                                                                                                            Similarity
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Tang L,
                                                                                                                                                                          BP;
                                                                                       Conservative
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L, Shahabuddin
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                                                                                       Gaps
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RESULT 684 ADJ59867

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RESULT 685
ADJ59775
ID ADJ597
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AC ADJ597
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DT 06-MAY
                                                                                                                                                                                                                                                                                         The present invention relates to an oligonuclectide anti-sense to e.g., cc initiation codon, coding region with 2-10 nucleotides of 5'-end and 3'-cc end of nucleic acid target comprising gene(s) chosen from e.g. cc interleukin (II)-4 receptor, II-5 receptor or salts of the cligonuclectide and optionally surfactant operatively linked to the cc oligonuclectide. The method is useful for preventing or treating a cc respiratory or lung disease, which involves administering to the airways of a subject an effective amount of an inhibitor. The oligonuclectide is cc of a subject an effective amount of an inhibitor. The oligonuclectide is cc useful for production of a medicament for the prevention and/or treatment of a respiratory or lung disease is chosen from airway inflammation, allergy(ies), asthma, impeded cc respiration, cystic fibrosis (CF), chronic obstructive pulmonary diseases (CODD), allergic rhinitis (AR), acute respiratory distress syndrome cc (ARDS), pulmonary hypertension, lung inflammation, bronchitis, airway construction. The present sequence represents an oligonuclectide of the
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Best Local S
Matches 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes (CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Shahabuddin
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                                                                                                                                                                                                                                                    Sequence 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      interleukin; IL-4 receptor; IL-5 receptor; lung disease;
   06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (EPIG-) EPIGENESIS PHARM INC.
                                                           ADJ59775
                                                                                                                                                              638
                                                                                                                                                                                             l Similarity
19; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        e.g., asthma.
                                                                                                                                   TGTCACCCAGGCTGGAGTG 656
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tang
lin S,
                                                           standard;
                                                                                                                                                                                                                                                       BP; 3
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 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sandrasagra
H, Cong H;
                                                             DNA;
                                                                                                                                                                                     1.9%; 5c.
100.0%; Pr
                                                                                                                                                                                                                                                     5 C; 7 G; 5 T; 0 U; 0 Other;
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                                                                                                                                                                                                           Score 19;
Pred. No.
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                                                                                                                                                                                             Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         impeded respiration;
                                                                                                                                                                                                                           DB 1;
                                                                                                                                                                                                           1.2e+03;
                                                                                                                                                                                                                           Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Miller
                                                                                                                                                                                              Indels
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RESULT 686
ADM14845/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CC The present invention relates to an oligonucleotide anti-sense to e.g., cc initiation codon, coding region with 2-10 nucleotides of 5'-end and 3'-cc end of nucleic acid target comprising gene(s) chosen from e.g. cc interleukin (IL)-4 receptor, IL-5 receptor or salts of the cc oligonucleotide and optionally surfactant operatively linked to the cligonucleotide. The method is useful for preventing or treating a cc respiratory or lung disease, which involves administering to the airways of a subject an effective amount of an inhibitor. The oligonucleotide is useful for production of a medicament for the prevention and/or treatment cc fa respiratory or lung disease. The respiratory or lung disease is cc hosen from airway inflammation, allergy(ies), asthma, impeded cc respiration, cystic fibrosis (CF), chronic obstructive pulmonary diseases (COPD), allergic rhinitis (AR), acute respiratory distress syndrome (CORDS), pulmonary hypertension, lung inflammation, bronchitis, airway inflammation, allergy (ies), asthma in the content of the content o
                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g., asthma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     interleukin; IL-4 receptor; IL-5 receptor; lung disease;
airway inflammation; allergy; asthma; impeded respiration;
cystic fibrosis; acute respiratory distress syndrome;
pulmonary hypertension; lung inflammation; bronchitis; oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Oligonucleotide associated to RANTES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2004-203534/19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               29-JUL-2002; 2002US-0399076P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20
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                                                                    Human mPGES-1
                                                                                                                      01-JUL-2004
                                                                                                                                                                                                         ADM14845 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      obstruction. The present sequence represents an oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (EPIG-)
                                                                                                                                                                                                                                                                                                                                                                728 GAGTAGCTGGGACTACAGG 746
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2; SEQ ID NO 631; 85pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                             19;
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                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                      (first entry)
                                                                    chimeric antisense oligonucleotide SEQ ID NO:1032.
                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                             1.2e+03;
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chimeric; antisense oligonucleotide; phosphorothioate; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1

human; inhibitor;

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The present sequence represents a chimeric antisense oligonucleotide CC targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The CC human mpGES-1 gene is located on chromosome 9, more specifically to 9934.3. The present invention also describes: (1) antisense compounds, CC having a sequence comprising 8-30 bp targeted to a nucleic acid encoding CC mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and CC inhibits its expression; (2) a method of inhibiting the expression of CC mpGES-1 in cells or tissues; and (3) a method of treating an animal CC having a disease or condition associated with mpGES-1 mpGES-1 chimeric CC antisense oligonucleotides and antisense compounds have cytostatic; CC antidiabetic, immunomodulator, cardiant, neuroprotective, antiarthritic, vasotropic, CC ophthalmological, immunomodulatory and cardiovascular activities, and can be used for preparing a composition for treating a disease or CC condition associated with mpGES-1 e.g., inflammation, Alzheimer's CC condition associated with mpGES-1 e.g., inflammation, Alzheimer's condition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's condition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's condition for treating a disease or condition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's condition for treating a disease or con
                                                                              Query Match
Best Local S
Matches 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g. inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
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                                                                                                                                                                                                                                                                                                                                ophthalmic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (PHAA ) PHARMACIA CORP
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846 GCCTCGGCCTCCCAAAGTG 864
                                                                              l Similarity
19; Conserv
                                                                                                                                                                                                                                              BP; 3
                                                                                                                                                                                                                                                                                                                                immunological,
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16. .20
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residues are 5-methylcytidines"
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/mod_base= OTHER
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                                                                                                                     100.0%;
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Pred. No.
                                                                                  Mismatches
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antisense oligonucleotides and antisense compounds have cytostatic

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RESULT 687
ADM14508/c
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targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9934.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric
                                                                                                                                                                                                                                                                                                                                                                                                                                          New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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                                                                                                                                                                                                                                                                                                                                                Claim 4; SEQ ID NO 695; 132pp; English.
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residues are 5-methylcytidines"
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     New antisense compound,
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16. .20
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     having a sequence targeted to a nucleic
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Pred. No.
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Homo sapiens. Synthetic.

modified_base

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mod_base= OTHER

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16. .20 /*tag=

note= "2'-0-methocyethyls"

note= "2'-0-methoxyethyls'

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/note "phosphorothicate linkages and all cytidine residues are 5-methylcytidines"

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RESULT 689
ADM15012/c
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Best Local S
Matches 19
                                                chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mpcES-1; mpcES-1 inhibitor; microsomal prostaglandin E2 synthase; mpcES-1; mticric; antidiabetic; microsomal prostaglandin E2 synthase; inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder;
                                                                                                                                                                                                                                                                   01-JUL-2004
                                                                                                                                                                                                                                                                                                                                              ADM15357 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 4; SEQ ID NO 1199; 132pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., encoding mPGES-1, useful for preparing a composition for treating e.g., arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2004-305094/28
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                                                                                                                                                                                                                               mPGES-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GCCTCCTGAGTAGCTGGGA 739
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 5 A; 7 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                             chimeric antisense oligonucleotide SEQ ID NO:1544.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%;
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Pred. No.
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RESULT 691 ADM15184/c

ADM15184

standard; DNA;

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                                                                                              Matches
                                                                                                                                     Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostagiandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9q34.3. The present invention also describes: (1) antisense compounds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.ginflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                        ophthalmic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 4; SEQ ID NO 1544; 132pp;
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19
                                                                                           l Similarity
19; Conserv
                           TTTTTGTATTTTTAGTAGA 787
                                                                                                                                                                                         20 BP; 12 A; 3 C; 0 G; 5 T; 0 U; 0 Other;
  TTTTGTATTTTAGTAGA 1
                                                                                                                                                                                                                                      immunological,
                                                                                              Conservative
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                                                                                                               1.9%;
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                                                                                                                                                                                                                                        cardiovascular
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    antisense compounds,

                                                                                                                                        Length
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                                                                                              Gaps
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microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotectie; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; hlzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder;
                                antidiabetic, immunomodulator, cardiant, neuroprotective, asotropic, antidiabetic, immunomodulatory nootropic, antiarthritic, vasotropic, antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's condition associated with mpGES-1 e.g., inflammation, and condition associated with mpGES-1 e.g., inflamm
                                                                                                                                                                                                                   targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9334.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibite its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human mPGES-1 chimeric antisense oligonucleotide SEQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 4;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQ ID NO 1371; 132pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             sequence represents a
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disease, arthritis, diabetes, cancer, iscl ophthalmic, immunological, cardiovascular

The invention relates to oligonucleotides anti-sense to an initiation codon, coding region, 5' or 3' intron-exon junction, intron or region with 2-10 nucleotides of the 5'-end or 3'-end of a nucleic acid target chosen from a gene encoding interleukin (II)-4 receptor, interleukin (II)-5 receptor, CCR1, CCR3, Ectaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, tryptase b, PDE4 A, PDE4 B, PDE4 C or PDE4 D. The invention also relates to a method of screening a candidate compound that binds to one or more nucleic acid target(s) or expressed product(s), for the prevention and/or treatment of a respiratory or lung disease. The oligonucleotides are useful for reducing or inhibiting expression of a

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RESULT 692
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Best Local S
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                                                                                                                                                                                                                                                                                                                                           Novel single or multiple target oligonuclectide anti-sense to e.g. Cinitiation codon, intron of respiratory disease-relevant gene e.g. CC RANTES, MCP4, useful for prophylaxis or treating respiratory disease
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23-APR-2002; 2002WO-US013143.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human oligonucleotide #631.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADO45265
                                                                                                                                                                                                                                                                Claim 2; SEQ ID NO 631; 174pp;
                                                                                                                                                                                                                                                                                                                                      RANTES, MCP4, useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Shahabuddin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (AGUI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (TANG/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (NYCE/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (SAND/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           inflammation; bronchitis; airway obstruction; bronchoconstriction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     771
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TANG L.
AGUILAR D.
MILLER S.
SHAHABUDDIN
LU H.
CONG H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NYCE J W
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SANDRASAGRA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TTTGTATTTTAGTAGAGA 789
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sandrasagra
in S, Lu H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         တ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Þ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                A A, Cong
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1.9%;
                                                                                                                                                                                                                                                                                                                                      for prophylaxis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ₽P
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ŗ
                                                                                                                                                                                                                                                                   English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Aguilar D,
                                                                                                                                                                                                                                                                                                                                        or treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1;
. 1.2e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Miller
                                                                                                                                                                                                                                                                                                                                        respiratory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                    CCR1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GF.
                                                                                                                                                                                                                                                                                                                                           e.9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0
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RESULT 693
ADO45357
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CCR1, CCR3, Botaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, tryptase b, DD24 A, DD24 B, DD24 C, or DD24 D. The oligonucleotides are useful for preventing or treating a respiratory or lung disease. The respiratory or lung disease is associated with hyper-responsiveness to and/or increased levels of, adenosine and/or levels of adenosine A receptor(s), and/or asthma and/or lung allergies associated with respiratory or lung disease inflammation or an inflammatory disease. The respiratory or lung disease is chosen from airway inflammation, allergy, asthma, impeded respiration, cystic fibrosis (CP), chronic obstructive pulmonary disease (CPD) allergic rhinitis, acute respiratory distress syndrome, pulmonary hypertension, lung inflammation, bronchitis, airway obstruction or the content of t
                                                                                                                                                                                                                                                                                            (SAND/) SANDRASAGRA F
(TANG/) TANG L.
(AGUI/) AGUILAR D.
(MILL/) MILLER S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; 88; interleukin-4 receptor; IL-4; interleukin-5 receptor; IL-5; CCR1; CCR3; Botaxin-1; RANTES; MCP4; CD23; ICAN; VCAM; tryptase a; tryptase b; PDE4 A; PDE8 B; PDE4 C; PDE4 D; respiratory disease; lung disease; hyper-responsiveness; adenosine; adenosine A receptor; asthma; lung allergy; inflammation; inflammatory disease; althma; lung allergy; inflammation; inflammatory disease; airway inflammation; allergy; impeded respiration; cystic fibrosis; CF chronic obstructive pulmonary disease; COPD; allergic rhinitis; acute respiratory distress syndrome; pulmonary hypertension; acute respiratory distress syndrome; pulmonary hypertension;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AD045357
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 5 A; 4 C; 8 G; 3 T; 0 U; 0 Other;
Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codon, intron of respiratory disease-relevant gene e.g.
                                                                                WPI; 2004-293804/27.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human oligonucleotide #723
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADO45357 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          bronchoconstriction. This sequence represents an oligonucleotide of the
                                                                                                                                Nyce JW, Sandrasagra
Shahabuddin S, Lu H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                          23-APR-2002; 2002WO-US013135.
23-APR-2002; 2002WO-US013143.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US2004049022-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        25-JUL-2003; 2003US-00627930.
                                                                                                                                                                                                                                                                                                                                                                                                        (NYCE/) NYCE J W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            or mRNA encoding interleukin-4 receptor, interleukin-5 receptor,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         728 GAGTAGCTGGGACTACAGG 746
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              N
                                                                                                                                                                                                                LU H.
                                                                                                                                                                                                                                                                     SHAHABUDDIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GAGTAGCTGGGACTACAGG 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.9%; Score 19; DB ilarity 100.0%; Pred. No. 1.2 Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         bronchitis; airway obstruction; bronchoconstriction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA;
                                                                                                            d A,
Cong
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20
                                                                                                                                  Tang
mg H;
                                                                                                                                                           Ļ
                                                                                                                                                         Aguilar D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1; Lo
1.2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 20;
                                                                                                                                                              Miller
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ç
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RESULT 694
ADPO8716
ID ADPO87
XX ADPO87
AC ADPO87
XX Z6-AUG
XX Z6-AUG
XX Dreast
XW GP6; C
XW Single
XX W02004
XX C5-NON
XX C5-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    밁
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CC chosen from a gene encoding interleukin (II)-4 receptor, interleukin (II)

CC -5 receptor, CCR1, CCR3, Botaxin-1, RANTES, MCP4, CD23, ICAM, VCAM,

CC tryptase a, tryptase b, PDE4 A, PDE4 B, PDE4 C. TO PDE4 D. The invention

CC also relates to a method of screening a candidate compound that binds to

CC one or more nucleic acid target(s) or expressed product(s), for the

CC prevention and/or treatment of a respiratory or lung disease. The

CC oligonucleotides are useful for reducing or inhibiting expression of a

CC gene or mRNA encoding interleukin-4 receptor, interleukin-5 receptor,

CC CCR1, CCR3, Botaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a,

CC tryptase b, PDE4 A, PDE4 B, PDE4 C, or PDE4 D. The oligonucleotides are

CC tryptase b, PDE4 A, PDE4 B, PDE4 C, or PDE4 D. The oligonucleotides are

CC tryptase b, and/or lasting a respiratory or lung disease. The

CC respiratory or lung disease is associated with hyper-responsiveness to

CC and/or increased levels of, adenosine and/or levels of adenosine A

CC receptor(s), and/or asthma and/or lung allergies associated with

CC inflammation or an inflammation, allergy, asthma, impeded respiration,

CC setic fibrosis (CF), chronic obstructive pulmonary disease (COPD),

CC oligonucleotide of the

CC bronchoconstriction. This sequence represents an oligonucleotide of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              breast cancer; cytostatic; gene therapy; human; platelet glycoprotein VI; GP6; GPIV; GPVI; chromosome 19q13.4; ss; PCR; primer; SNP; single nucleotide polymorphism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Extend primer 53 used to genotype human glycoprotein VI polymorphism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 with 2-10 nucleotides of the 5'-end or 3'-end of a nucleic acid target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to oligonucleotides anti-sense to an initiation codon, coding region, 5' or 3' intron-exon junction, intron or region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     asthma.
                                                                                                                                   Roth RB,
                                                                                                                                                                                                                                                             25-NOV-2002;
24-JUL-2003;
                                                                                                                                                                                                                                                                                                                                                               25-NOV-2003; 2003WO-US037966
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO2004047767-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         26-AUG-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADP08716 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                           10-JUN-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 638 TGTCACCCAGGCTGGAGTG 656
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1.9%; Score 19;
1 Similarity 100.0%; Pred. No.
19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        N
                                                                                                                                                                                             SEQUENOM INC
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                                                                                                                                Nelson MR,
                                                                                                                                                                                                                                                             2002US-0429136P.
2003US-0490234P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ID NO 723; 174pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; 5 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                                   Braun
                                                                                                                                   Þ
                                                                                                                                   Kammerer SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1.2e+03;
hes 0;
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                                                                                                                                Reneland
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 20;
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WPI; 2004-441082/41.

Identifying a subject at risk of breast cancer by detecting the presence

or absence diagnosing,

preventing

of one or more nucleotide polymorphic variations, useful for

and/or treating breast

cancer.

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FFXEXDOCOCOCOCOC
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Best Local S
Matches 19
                                                                                                                                                                                                       20-FEB-2002;
11-MAR-2002;
06-JUN-2002;
29-AUG-2002;
05-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to a novel method for identifying a subject at risk of breast cancer which comprises detecting the presence or absence of one or more polymorphic variations associated with breast cancer in a nucleic acid sample from a subject. The method of the invention has cytostatic applications and may be useful for identifying a risk of breast cancer,
                                                                                                                                                                                                                                                                                                                                                                                                     antiarteriosclerotic; neuroprotective; nootropic; antiparkinsonian; anticonvulsant; pulmonary disease; restenosis; atherosclerosis; Alzheimer's, Parkinson's, epilepsy; dementia; huntington's, amyotrophic lateral sclerosis; gene therapy; ss; DNA-RNA hybrid; PKR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADG30202 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                    20-FEB-2003;
                                                                                                                                                                                                                                                                                                               12-SEP-2003
                                                                                                                                                                                                                                                                                                                                        WO2003074654-A2
                                                                                                                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                        double-stranded short interfering nucleic acid; siNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PKR-targeted sina DNA-RNA hybrid -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             26-FEB-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADG30202;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           nucleotide
                                    New double-stranded short interfering nucleic acid molecule, useful for down-regulating the expression of an endogenous mammalian target gene of treating diseases that respond to modulation of gene expression or
                                                                                                                           Mcswiggen
                                                                                                                                                      (SIRN-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            well as therapeutic and prophylactic treatments that specifically reget breast cancer, such as gene therapy. The current sequence is that an Extend primer of the invention which was used to genotype single cleotide polymorphisms within human glycoprotein VI (platelet) (GP6; IV,GPVI) DNA which is located at chromosomal position 19q13.4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         719
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19; Conserv
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                                                                                                                              J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Page 83;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP; 3 A; 6 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                             2002US-0358580P.
2002US-0363124P.
2002US-0366782P.
2002US-0406784P.
2002US-0408378P.
2002US-0409293P.
2003US-0440129P.
                                                                                                                Usman
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                                                                                                                                                                                                                                                                                       2003WO-US005028
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry
                                                                                                               Beigelman L, Cho
aman N, Thompson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         286pp; English
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Pred. No.
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                                                                                                                               Pavco
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                                                                                                                              'n
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
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                                                      or
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useful for down-regulating the expression of an endogenous mammalian target gene and therefore in the treatment of any disease or condition that responds to modulation of gene expression or activity in a cell, tissue or organism. The disease or condition may include pulmonary disease such as restenosis, atherosclerosis, Alzheimer's disease, parkinson's disease, epilepsy, dementia, huntington's disease or amyotrophic lateral sclerosis. Purthermore, the siNA may be utilised for gene therapy applications. The current sequence is that of the siNA DNA-RNA hybrid of the invention.
                                                                                                                                                                                                                                                                                                     The invention relates to a double-stranded short interfering nucleic ((6iNA) molecule that down-regulates expression of an endogenous mamma trarget gene comprising one or more chemical modifications and each stof the double-stranded siNA comprises about 21 nucleotides. The siNA the invention demonstrates antiarteriosclerotic, neuroprotective, nootropic, antiparkinsonian and anticonvulsant activities and may be
                                                                                                                                                                                                                                                                                                                                                                                                                           interfering nucleic acid an endogenous mammalian
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Best Loc
Matches
                                                 Query Match
                                 Local Similarity
nes 19; Conserv
               1117 GGTCTCAAACTCCTGACCT 1135
19
GGTCTCAAACTCCTGACCT 1
                                  Conservative
                                          1.9%;
                                  0;
                                           Score 19;
Pred. No.
                                   Mismatches
                                             T BB
                                            .3e+03
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                                    Indels
                                    0;
                                    Gaps
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Sequence 21 BP; 5 A; 3 C; 7 G; 2 T; 4 U;

0 Other;

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RESULT 696
                                    20-MAY-2004
                                                                       ADL25334;
 Intestinal
                                                                                                         ADL25334
                                                                                                        standard; DNA; 21 BP
epithelium/peyer's patch M cell-associated PCR primer #479.
                                    (first entry)
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ID ADIL25334/c
ID ADIL253
XX
AC ADIL253
XX
XX ADIL253
XX
XX Intest
XX Intest autoimmune disease; haemolytic anaemia; rheumatoid art Grave's disease; multiple sclerosis; allergy; asthma; immune system disorder; hypersensitivity; anaphylaxis; blood group incompatibility; ss; PCR; primer. inflammatory ntestinal epithelium cell development; bowel disease; glutenenteropathy; infectious disease; rheumatoid arthritis; (lergy; asthma; diabetic peyer's patch M cell development, dermatitis;

Macaca fascicularis

WO200280852-A2

17-OCT-2002.

04-APR-2002; 2002WO-US010873

04-APR-2001; 2001US-0281416P

(DIGI-) DIGITAL GENE TECHNOLOGIES

Brayden Ď, Byrne D, O'mahony DJ, Evans CF, Mah SP, ۲

日

WPI; 2003-075470/07

Novel isolated or purified polypeptide encoded by genes associated with intestinal epithelium or M cell development, differentiation or function useful for treating autoimmune diseases and infectious diseases. function

Disclosure; SEQ ID NO 844; 152pp; English.

The invention comprises DNA sequences which are associated with intestinal epithelium and peyer's patch M cells. The DNA sequences of the invention are useful for assessing, modifying, modulating or regulating intestinal epithelium or M cell development. The DNA sequences of the invention are also useful in the treatment of: inflammatory bowel glutenenteropathy, of the

Example 24;

QES

ID NO 768;

593pp; English

. haemolytic anaemia, rheumatoid arthritis, dermatitis, Grave's

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RESULT 697
AAZ25166
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                                            The present invention describes a restriction primer for eukaryotic short interspersed repeated sequences (SINE), which has one or more additional bases that are a mismatch to, or are unrelated to, the 3'-terminal end of the SINE. The annealing temperature of the primer to the DNA sequence is kept higher than the fusion temperature of the primer during polymerase chain reaction (PCR). The PCR fragments obtained are subjected to electrophoresis to obtain a fingerprint. By comparing the polymorphs from the electrophoresis band pattern, eukaryotic individuals are city of the primer is used for amplifying a eukaryotic descriptionucleic acid (DNA) sequence, pinched between SINE sequences by colymerase chain reaction (PCR) fingerprinting. In particular it may be used individual identification of humans for medical and legal capplications and ecological studies. DNA specimens in traces applications and ecological studies. DNA specimens in traces chain reaction (PCR).

CC approximately 10 ng in mass) can be used for individual discrimination of eukaryotes using the primer in a polymerase chain reaction (PCR).

CC ALZ25143 to ALZ25191 represent specifically claimed examples of primers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              disease, multiple sclerosis, allergy, asthma and diabetic mellitus), diseases or disorders of the immune system, hypersensitivity, anaphylaxis, and blood group incompatibility. The present DNA sequence represents a PCR primer that was used to amplify an intestinal epithelium/peyer's patch M cell-associated DNA sequence of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; short interspersed repetitive element; SINE; PCR; primer; Oncorhynchus; restriction primer; short interspersed repeated sequence; eukaryote, restricted polymerase chain reaction fingerprinting; identification; DNA specimen; discrimination; ss.
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                                                                                                                                                                                                                                                                                                                                                                              Claim 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                 Restriction primer for distinguishing individuals with short interspersed repeated sequence of eukaryotes by restricted polymerase chain reaction
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TTCTCCTGTCTCAGCCTCC 3
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ilarity 100.0%;
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Sequence 22 BP; 5

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0 U; 0 Other

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GATTACAGGCGTGAGCCAC

887

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Mismatches

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GATTACAGGCGTGAGCCAC

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Best Local S
Matches 19
Query Match
Best Local Similarity
Matches 19; Conserv
                                                                                                                                                                                          The present invention describes a restriction primer for eukaryotic short interspersed repeated sequences (SINB), which has one or more additional bases that are a mismatch to, or are unrelated to, the 3'-terminal end of the SINB. The annealing temperature of the primer to the DNA sequence is kept higher than the fusion temperature of the primer during polymerase chain reaction (PCR). The PCR fragments obtained are subjected to electrophoresis to obtain a fingerprint. By comparing the polymorphs from the electrophoresis band pattern, eukaryotic individuals are distinguished. The primer is used for amplifying a eukaryotic deoxyribonucleic acid (DNA) sequence, pinched between SINB sequences by polymerase chain reaction (PCR) fingerprinting. In particular it may be used individual identification of humans for medical and legal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human, short interspersed repetitive element; SINE; PCR; primer; Oncorhynchus; restriction primer; short interspersed repeated sequence; eukaryote; restricted polymerase chain reaction fingerprinting; identification; DNA specimen; discrimination; 88.
                                                                                                                        applications and ecological studies. DNA specimens in traces (approximately 10 ng in mass) can be used for individual discrimination of eukaryotes using the primer in a polymerase chain reaction (PCR).

AAZ25143 to AAZ25191 represent specifically claimed examples of primers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Restriction primer for distinguishing individuals with short interspersed repeated sequence of eukaryotes by restricted polymerase chain reaction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic
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                                                                     Sequence 22
                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1999-583348/50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        10-JUL-1998;
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                                                                                                        present invention
                                                                                                                                                                                                                                                                                                                                                                                                                            Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1.9%;
llarity 100.0%;
Conservative (
                                                                       BP; 7 A; 5 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        98JP-00195692.
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                    100.0%;
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1.3e+03;
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RESULT 700
AAZ25170
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                                                                                                                                                                                                                                                                                The present invention describes a restriction primer for eukaryotic short contemporared repeated sequences (SINE), which has one or more additional the bases that are a mismatch to, or are unrelated to, the 3'-terminal end of the SINE. The annealing temperature of the primer to the DNA sequence is comparing the primer to the DNA sequence is comparing the polymorphs from the electrophoresis to obtain a fingerprint. By comparing the polymorphs from the electrophoresis band pattern, eukaryotic individuals are the electrophoresis band pattern, eukaryotic individuals are constituted in the primer is used for amplifying a eukaryotic deoxyribonucleic acid (DNA) sequence, pinched between SINE sequences by colymerase chain reaction (PCR) fingerprinting. In particular it may be used individual identification of humans for medical and legal applications and ecological studies. DNA specimens in traces applications and ecological studies. DNA specimens in traces chain reaction (PCR).

ANAZ25143 to ANAZ25191 represent specifically claimed examples of primers
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Restriction primer for distinguishing individuals with short interspersed repeated sequence of eukaryotes by restricted polymerase chain reaction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10-JUL-1998;
                                                                                                                                                                                                                                        Sequence 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; short interspersed repetitive element; SINE; PCR; primer
              AAZ25170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (NORQ ) NORINSUISANSHO SUISANCHO YOSHOKU KENKYUSHOCHO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                              AAZ25170 standard; DNA; 22
                                                                                                                                                                                                                                                                          the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1999-583348/50.
                                                                                                                                                    869
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                                                                                                                                                                                l Similarity
19; Conserv
                                                                                                                                     GATTACAGGCGTGAGCCAC 887
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Page 3; 17pp; Japanese.
                                                                                                                                                                                                                                                                          present
                                                                                                                       GATTACAGGCGTGAGCCAC 19
                                                                                                                                                                                                                                          BP; 6 A; 5 C; 7 G; 4 T; 0 U; 0 Other;
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                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             98JP-00195692
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                                                                                                                                                                                                                                                                        invention
                                                                                                                                                                          1.9%; 5r/
100.0%; Pr/
0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP.
                                               ВP
                                                                                                                                                                                               Score 19;
Pred. No.
                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                DB 1;
                                                                                                                                                                                               1.3e+03;
                                                                                                                                                                                  0,
                                                                                                                                                                                                             Length 22;
                                                                                                                                                                                  Indels
                                                                                                                                                                                  0
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RESULT 701
AAZ25172
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                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human short interspersed repetitive element \mathbf{d}
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; short interspersed repetitive element; SINE; PCR; primer; Oncorhynchus; restriction primer; short interspersed repeated sequence; eukaryote; restricted polymerase chain reaction fingerprinting; identification; DNA specimen; discrimination; ss.
Human; short interspersed repetitive element; SINE; PCR; primer; Oncorhynchus; restriction primer; short interspersed repeated seeukaryote; restricted polymerase chain reaction fingerprinting; identification; DNA specimen; discrimination; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Restriction primer for distinguishing individuals with short interspersed repeated sequence of eukaryotes by restricted polymerase chain reaction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28-JUN-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 22 BP; 5 A; 5 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     of eukaryotes using the primer in a polymerase chain reaction (PCR).
AAZ25143 to AAZ25191 represent specifically claimed examples of primers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   fingerprinting.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               10-JUL-1998;
                                                                                         Human short interspersed repetitive element PCR primer #30
                                                                                                                              13-DEC-1999
                                                                                                                                                              AAZ25172;
                                                                                                                                                                                                 AAZ25172 standard; DNA; 22 BP
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                                                                                                                                                                                                                                                                                                                                                                                 Local
                                                                                                                                                                                                                                                                                                                           869 GATTACAGGCGTGAGCCAC 887
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  6
                                                                                                                                                                                                                                                                                                                                                            19;
                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                         GATTACAGGCGTGAGCCAC 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       present invention
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                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                              (first entry)
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                                                                                                                                                                                                                                                                                                                                                                               1.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SUISANCHO YOSHOKU KENKYUSHOCHO
                                                                                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                                                                                                                                                             Score 19; DB 1; Length 22; Pred. No. 1.3e+03;
                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           primer #28
                                                                                                                                                                                                                                                                                                                                                                Indels
                                          repeated sequence;
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AAZZ5159
ID AAZZ
XX
AC AAZZ
XX
DT 13-[
XX
DB Huma
XX
KW Huma
KW Onco
KW euk
KW ide
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KW 129
DS Hom
XX
PD JP2
XX
PD 28-
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                                                                                                                                                                                                                                                                                                                                                                   RESULT 702
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention describes a restriction primer for eukaryotic short interspersed repeated sequences (SINE), which has one or more additional bases that are a mismatch to, or are unrelated to, the 3' terminal end of the SINE. The annealing temperature of the primer to the DNA sequence is kept higher than the fusion temperature of the primer during polymerase chain reaction (PCR). The PCR fragments obtained are subjected to electrophoresis to obtain a fingerprint. By comparing the polymorphs from the electrophoresis band pattern, eukaryotic individuals are distinguished. The primer is used for amplifying a eukaryotic deoxyribonucleic acid (DNA) sequence, pinched between SINE sequences by polymerase chain reaction (PCR) fingerprinting. In particular it may be used individual identification of humans for medical and legal applications and ecological studies. DNA specimens in traces (approximately 10 ng in mass) can be used for individual discrimination of the primer in a polymerase chain reaction (DCP)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 6; Page 4; 17pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Restriction primer for distinguishing individuals with short interspersed repeated sequence of eukaryotes by restricted polymerase chain reaction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1999-583348/50.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10-JUL-1998;
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                   28-JUN-1999
                                                    JP2913035-B1
                                                                                                                                   Oncorhynchus; restriction primer; short interspersed repeated sequence; eukaryote; restricted polymerase chain reaction fingerprinting; identification; DNA specimen; discrimination; ss.
                                                                                                                                                                                                                                                                                               AAZ25159
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               fingerprinting.
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                                                                                                     Synthetic
                                                                                                                                                                                          Human; short interspersed repetitive element; SINE; PCR; primer;
                                                                                                                                                                                                                          Human short interspersed repetitive element PCR primer #17.
                                                                                                                                                                                                                                                              13-DEC-1999
                                                                                                                                                                                                                                                                                                                                 AAZ25159 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (NORQ ) NORINSUISANSHO SUISANCHO YOSHOKU KENKYUSHOCHO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                eukaryotes using the primer in a polymerase chain reaction (PCR).
225143 to AAZ25191 represent specifically claimed examples of primers
                                                                                    sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                      869 GATTACAGGCGTGAGCCAC 887
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19;
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                                                                                                                                                                                                                                                                                                                                                                                                                      GATTACAGGCGTGAGCCAC 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP; 6 A; 6 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                              (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 invention
                                                                                                                                                                                                                                                                                                                                 DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1.9%;
                                                                                                                                                                                                                                                                                                                                   22
                                                                                                                                                                                                                                                                                                                                   ВP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1; Length 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1.3e+03;
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RESULT 703
AAZ25163
ID AAZ251
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CC ALRZ5143 to ARZ55191 represent specifically claimed examples of primers from the pracet in a polymerase chain reaction (PCR).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
Matches 19; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Restriction primer for distinguishing individuals with short interspersed repeated sequence of eukaryotes by restricted polymerase chain reaction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10-JUL-1998;
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                                                                                                                                                                                                                                 Synthetic.
                                                                                                                                                                                                                                                                  eukaryote; resti
identification;
                                                                                                                                                                                                                                                                                Oncorhynchus; restriction primer; short interspersed repeated sequence; enkaryote; restricted polymerase chain reaction fingerprinting;
                                                                                                                                                                                                                                                                                                                  Human; short interspersed repetitive element; SINE; PCR; primer;
                                                                                                                                                                                                                                                                                                                                                     Human short interspersed repetitive element PCR primer #21.
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                                                                                                                                                                                                                                                                                                                                                                                                                       AAZ25163;
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             WPI; 1999-583348/50
                                                                                10-JUL-1998;
                                                                                                               10-JUL-1998;
                                                                                                                                                 28-JUN-1999
                                                                                                                                                                                                                   Homo
                                               (NORQ ) NORINSUISANSHO SUISANCHO YOSHOKU KENKYUSHOCHO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               rom the
                                                                                                                                                                                                                 sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GATTACAGGCGTGAGCCAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             present invention
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                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
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                                                                                98JP-00195692.
                                                                                                               98JP-00195692
                                                                                                                                                                                                                                                                    DNA specimen; discrimination;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1.9%;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 22
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RESULT 704
AAZ25169
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               IJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   fingerprinting.
                                                                                                                                                                                                                                                                                                                                                                             Human; short interspersed repetitive element; SINE; PCR; primer; Oncorhynchus; restriction primer; short interspersed repeated sequence; eukaryote; restricted polymerase chain reaction fingerprinting;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    repeated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Restriction primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAZ25169 standard; DNA; 22
                                                                                      Restriction primer for distinguishing individuals with short interspersed repeated sequence of eukaryotes by restricted polymerase chain reaction
                                                                                                                                                                                                                                                                                      JP2913035-B1
                                                                                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                          Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      13-DEC-1999
                                                                                                                                    WPI; 1999-583348/50
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                                                                                                                                                                                                                            10-JUL-1998;
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                                                                                                                                                                                                                                                                                                                                                                  identification;
                                                                                                                                                              (NORQ ) NORINSUISANSHO SUISANCHO YOSHOKU KENKYUSHOCHO
                                                                                                                                                                                                                                                                                                                 sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Page 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GATTACAGGCGTGAGCCAC 19
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Б
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   for distinguishing individuals with short interspersed of eukaryotes by restricted polymerase chain reaction
                                                                                                                                                                                                                                                                                                                                                                specimen;
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                                                                                                                                                                                                                                                                                                                                                                                                                                        repetitive element PCR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 19;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                  discrimination; ss.
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The present invention interspersed repeated

describes a restriction sequences (SINE), which

primer for eukaryotic short has one or more additional

The present invention describes a restriction primer for eukaryotic short interspersed repeated sequences (SINE), which has one or more additional bases that are a mismatch to, or are unrelated to, the 3'-terminal end of the SINE. The annealing temperature of the primer to the DAN sequence is kept higher than the fusion temperature of the primer during polymerase chain reaction (PCR). The PCR fragments obtained are subjected to electrophoresis to obtain a fingerprint. By comparing the polymorphs from the electrophoresis band pattern, eukaryotic individuals are distinguished. The primer is used for amplifying a eukaryotic deoxyribonucleic acid (DNA) sequence, pinched between SINE sequences by

Restriction primer for distinguishing individuals with short interspersed repeated sequence of eukaryotes by restricted polymerase chain reaction

Claim 6; Page 4; 17pp; Japanese.

6

Page

4; 17pp; Japanese

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RESULT 705
AAZ25171
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; short interspersed repetitive element; SINE; PCR; primer; Oncorhynchus; restriction primer; short interspersed repeated sequence; eukaryote; restricted polymerase chain reaction fingerprinting; identification; DNA specimen; discrimination; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           applications and ecological studies. DNA specimens in traces (approximately 10 ng in mass) can be used for individual discrimination of eukaryotes using the primer in a polymerase chain reaction (PCR).

AAZ25143 to AAZ25191 represent specifically claimed examples of primers
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                                                                                                                                                                                                                                                                        (NORQ ) NORINSUISANSHO SUISANCHO YOSHOKU KENKYUSHOCHO
                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
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RESULT 706
AAZ25160
The present invention describes a restriction primer for eukaryotic short clinterspersed repeated sequences (SINE), which has one or more additional bases that are a mismatch to, or are unrelated to, the 3'-terminal end of the SINE. The annealing temperature of the primer to the DNA sequence is kept higher than the fusion temperature of the primer during polymerase chain reaction (PCR). The PCR fragments obtained are subjected to electrophoresis to obtain a fingerprint. By comparing the polymorphs from the electrophoresis band pattern, eukaryotic individuals are distinguished. The primer is used for amplifying a eukaryotic deoxyribonucleic acid (DNA) sequence, pinched between SINE sequences by colymerase chain reaction (PCR) fingerprinting. In particular it may be used individual identification of humans for medical and legal cused individual identification of humans for medical and legal capplications and ecological studies. DNA specimens in traces (approximately 10 ng in mass) can be used for individual discrimination of eukaryotes using the primer in a polymerase chain reaction (PCR).

ANAZ25143 to ANZ25191 represent specifically claimed examples of primers
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                                                                                                                                                                                                                                                                                                                                                                               Claim 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                       Restriction primer for distinguishing individuals with short interspersed repeated sequence of eukaryotes by restricted polymerase chain reaction
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; short interspersed repetitive element; SINE; PCR; primer;
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                                        The present invention describes a restriction primer for eukaryotic short CC interspersed repeated sequences (SINE), which has one or more additional CC bases that are a mismatch to, or are unrelated to, the 3'-terminal end of CC the SINE. The annealing temperature of the primer to the DNA sequence is kept higher than the fusion temperature of the primer during polymerase CC chain reaction (PCR). The PCR fragments obtained are subjected to electrophoresis to obtain a fingerprint. By comparing the polymorphs from the electrophoresis band pattern, eukaryotic individuals are cdistinguished. The primer is used for amplifying a eukaryotic deoxyribonucleic acid (DNA) sequence, pinched between SINE sequences by CC deoxyribonucleic acid (DNA) sequence, pinched between SINE sequences by CC used individual identification of humans for medical and legal applications and ecological studies. DNA specimens in traces (approximately 10 ng in mass) can be used for individual discrimination of eukaryotes using the primer in a polymerase chain reaction (PCR).

ARZ25143 to ARZ25191 represent specifically claimed examples of primers
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                                                                                                                                                                                                                                                                                                         Claim
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ID ADG3
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AC ADG3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 708
                                                                                                                                      S
                                                                                                                                                                                                                                                                       The present invention describes a restriction primer for eukaryotic short clinterspersed repeated sequences (SINE), which has one or more additional CD bases that are a mismatch to, or are unrelated to, the 3'-terminal end of CD the SINE. The annealing temperature of the primer to the DNA sequence is contain reaction (PCR). The PCR fragments obtained are subjected to CD the interpretation (PCR). The PCR fragments obtained are subjected to CD the electrophoresis to obtain a fingerprint. By comparing the polymorphs from CD the electrophoresis band pattern, eukaryotic individuals are CD described. The primer is used for amplifying a eukaryotic cdeoxyribonucleic acid (DNA) sequence, pinched between SINE sequences by polymerase chain reaction (PCR) fingerprinting. In particular it may be used individual identification of humans for medical and legal CD applications and ecological studies. DNA specimens in traces applications are ecological studies. DNA specimens in traces constituted using the primer in a polymerase chain reaction (PCR). ARZES143 to ARZES191 represent specifically claimed examples of primers constituted to primers.
                                                                                                            뮍
                                                               RESULT 709
                                                                                                                                                                                     Query Match
Best Local (
                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Restriction primer for distinguishing individuals with short interspersed repeated sequence of eukaryotes by restricted polymerase chain reaction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        JP2913035-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; short interspersed repetitive element; SINE; PCR; primer; Oncorhynchus; restriction primer; short interspersed repeated sequence; eukaryote; restricted polymerase chain reaction fingerprinting; identification; DNA specimen; discrimination; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human short interspersed repetitive element PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAZ25165
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1999-583348/50.
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                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim
   ADG30198
                                  ADG30198 standard; RNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (NORQ ) NORINSUISANSHO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sapiens.
                                                                                                                                          698
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                                                                                                                                                                                     Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Page 4; 17pp; Japanese
                                                                                                                                          GATTACAGGCGTGAGCCAC 887
                                                                                                                                                                                                                                   22
                                                                                                            GATTACAGGCGTGAGCCAC 19
                                                                                                                                                                                                                                                                  present
                                                                                                                                                                                                                                   BP; 6 A;
                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             98JP-00195692.
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                                                                                                                                                                                                                                                                  invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA;
                                                                                                                                                                                     1.9%;
                                                                                                                                                                                                                                   6 C; 7 G; 3 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SUISANCHO YOSHOKU KENKYUSHOCHO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             22
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Pred. No.
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Matches
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11-MAR-2002; 2002US-0363124P.
06-JUN-2002; 2002US-03682P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0409293P.
15-JAN-2003; 2003US-0440129P.
                                                                                                                                                                                                                                          nootropic, antiparkinsonian and anticonvulsant activities and may be useful for down-regulating the expression of an endogenous mammalian that responds to modulation of gene expression or activity in a cell, tissue or organism. The disease or condition may include pulmonary diseases such as restenosis, atherosclerosis, Alzheimer's disease, parkinson's disease, epilepsy, dementia, huntington's disease or amyotrophic lateral sclerosis. Furthermore, the siNA may be utilised for gene therapy applications. The current sequence is that of the siNA DNA-RNA hybrid of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mcswiggen ,
Jamison S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   antiarteriosclerotic; neuroprotective; nootropic; antiparkinsonian; anticonvulsant; pulmonary disease; restenosis; atherosclerosis; alzheimer's; Parkinson's; epilepsy; dementia; huntington's; amyotrophic lateral sclerosis; gene therapy; ss; DNA-KNA hybrid; PKR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene comprising one or more chemical modifications and each strand of the double-stranded siNA comprises about 21 nucleotides. The siNA of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New double-stranded short interfering nucleic acid molecule, useful for down-regulating the expression of an endogenous mammalian target gene or for treating diseases that respond to modulation of gene expression or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           double-stranded short interfering
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PKR-targeted sinA DNA-RNA hybrid - SEQ ID 764
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       the invention demonstrates antiarteriosclerotic, neuroprotective,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       for treating activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-FEB-2003; 2003WO-US005028.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2003-731676/69.
                                                                                                                                                                                                  Sequence 23 BP; 4 A; 7 C; 3 G; 2
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ເ. Usman ທ
                                                                                                                          Similarity
                                               GGTCTCAAACTCCTGACCT 1135
                                                                                                  Conservative
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                                                                                                                       1.9%;
                                                                                                  5
                                                                                                                                                  Score 19;
                                                                                                  Pred. No. 1.40
5; Mismatches
20
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                                                                                                                                                                                                     U; 2 Other;
                                                                                                                          1.4e+03;
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                                                                                                                                                  Length 23;
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RESULT 710 AAT39493 ID AAT394 XX

AAT39493 standard; DNA; 22

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RESULT 711
AAX83018
ID AAX83018
XX AAX830
XX AAX830
XY 31-AUG
DT 31-AUG
XX Primer
XX Primer
XX Primer
XX Primer
XX Human;
KW Human;
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                                                                                                                                                                                                                                                                                                                           Query Match
Best Local
                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                             The present sequence is a PCR primer for exon 4 of the human steroidogenesis acute regulatory protein (hStAR) gene. The hStAR gene can be analysed for mutations to detect (e.g. prenatally) genetic defects associated with congenital lipoid adrenal hyperplasia (CAH), or its transmission to children. CAH can be treated by protein or gene replacement therapy, which can also be used to prevent or treat hypercholesterolaemia. A human adrenal cortex cDNA library was screened with a mouse StAR probe to isolate a 1.6 kb insert, including an ORF for a 285 residue protein. When it was cloned into pSPORT and expressed in COS-1 cells cotransfected with pP450scc abd pADX, it increased the level of pregnenolone synthesis from cholesterol or 20-alpha-hydroxycholesterol
                                          Human; WRN; Werner's syndrome; detection; diagnosis;
recessive disorder; phenotype; primer; RT-PCR; amplif
                                                                                                                    31-AUG-1999
                                                                                                                                                                                                                                                                                                                                                                        Sequence 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 36; 89pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Isolated human steroidogenesis acute regulatory protein gene - used detection of mutation(s) of this gene that cause congenital lipoid adrenal hyperplasia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Miller WL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         23-MAR-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         polymerase chain reaction; exon 4; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; steroidogenesis; acute regulatory protein; hStAR; analysis; mutation; detection; prenatal; genetic defect; congenital; protein; lipoid adrenal hyperplasia; treatment; prevention; gene; replacement therapy; hypercholesterolaemia; primer; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Steroidogenesis acute
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                                                                                                                                                                             AAX83018
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                                                                                                                                                                                                                                                                                863
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                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                             standard;
                                                                                                                                                                                                                                                                      TGCTGGGATTACAGGCGTGAGC 884
                                                                                                                                                                                                                                                     TGCTGGGATTATAGGCGTGAAC 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Lin D,
                                                                                                                                                                                                                                                                                                                                                                      BP; 5 A; 3 C; 8 G; 6 T; 0 U; 0 Other;
                                                                                       isolate
                                                                                                                                                                                                                                                                                                              Conservative
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PENNSYLVANIA.
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                                                                                                                                                                             DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Strauss
                                                                                     human WRN
                                                                                                                  entry)
                                                                                                                                                                                                                                                                                                                           90.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             regulatory protein exon 4 PCR primer Ex4S
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Pred. No. 1.
                                                                                     gene
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                                                                                       exons.
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                                            amplification; ss
                                                                                                                                                                                                                                                                                                                                        Length
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RESULT 712
AAC69375/c
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29-DEC-1995;
30-JAN-1996;
30-JAN-1996;
12-APR-1996;
         15-MAR-1999;
08-JUN-1999;
17-JUN-1999;
01-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Primers AAX83008-X83064 were used to RT-PCR amplify exons from the 5'3' ends of the human WRN gene (AAX83003) which encodes a protein relato Werner's syndrome. The products can be used for the detection and treatment of Werner's syndrome (WS), an autosomal recessive disorder a complex phenotype, as well as related diseases
                                                                                                                                                           X-linked adrenoleukodystrophy; cancer; gene therapy; genetic diagnosis;
prognosis; prophylaxis; drug screening; transgenic animal; ds.
                                                                                                                                                                                cerebrovascular disease; peripheral vascular disease; Alzheimer's disease; Niemann-Pick disease; Huntington's disease;
                                                                                                                                                                                                      Human ABC1 cholesterol transporter; chromosome 9q31;
ATP-binding cassette; HDL deficiency disorder; high density lipoprotein;
Tangier disease; TD; familial HDL deficiency; FHA; polymorphism;
cardiovascular disease; coronary artery disease; coronary restenosis;
                                                                                                                                                                                                                                                                                       29-JAN-2001
                                                                                                                                                                                                                                                                                                                                   AAC69375
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Isolated nucleic acid molecule encoding the WRN gene product - useful for detection and treatment of Werner's syndrome, and related diseases.
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                                                                                        21-SEP-2000
                                                                                                               WO200055318-A2
                                                                                                                                     Homo
                                                                                                                                                                                                                                                                 Human ABC1
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Similarity 90.9%;
                                                                                                                                                                                                                                                                                                                                    standard;
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                                                                                                                                                                                                                                                                BAC contig polymorphic site,
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                                                                   2000WO-IB000532
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95US-00580539.
96US-0010835P.
96US-00594242.
          99US-0124702P.
99US-0138048P.
99US-0139600P.
99US-0151977P.
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                                                                                                                                                                                                                                                                                                                                    DNA;
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Pred. No. 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Schellenberg
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                                                                                                                                                                                                                                                                 SEQ
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                                                                                                                                                                                                                                                                 NO:274
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CC The invention relates to the human ABC1 cholesterol transporter protein CC (B38082) and to nucleic acid sequences (C69120) which encode it. ABC1 is CC a member of the ATP-binding cassette (ABC transporter) superfamily of CC intracellular cholesterol trafficking in moncytes and fibroblasts, being CC involved in cholesterol efflux from the cell. The gene are associated control to the collection of the co
  TXAXAXBXBXXXXX
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                                                                                                                                                                                                                                                                     RESULT 713
                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hayden MR, Wilson AR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (UYBR-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2000-587528/55
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       biological activity, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example; Fig 11; 229pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
MKK4; mitogen activated protein kinase; MAPK; MAPK pathway; mutation; somatic; signal transduction; apoptosis; stress; cytokine; induction; phosphorylation; Jun kinase; JNK; p38; tumour; suppressor; loss of heterozygosity; LOH; cancer; detection; diagnosis; prognosis;
                                                                                                      Sequence tagged site (STS) 66I11.T7 probe forward PCR primer
                                                                                                                                               09-FEB-2000
                                                                                                                                                                                                                              AAZ32938 standard; DNA;
                                                                                                                                                                                                                                                                                                                                    22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   UNIV BRITISH COLUMBIA. XENON BIORESEARCH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                     20;
                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        polypeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        22
                                                                                                                                                                                                                                                                                                                                                                            TCCTCCTGCCTCAGCCTCCCAA 554
                                                                                                                                                                                                                                                                                                                                    TTCTCCTGCCTTAGCCTCCCAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 7 A;
                                                                                                                                               (first entry
                                                                                                                                                                                                                                                                                                                                                                                                                                         1.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     is useful for treating diseases associated with ABC1 e.g. Alzheimer's disease, "Huntington's disease and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2 C; 10 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pimstone
                                                                                                                                                                                                                                                                                                                                                                                                                     <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 18.8;
Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                           1.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 22;
                                                                                                                                                                                                                                                                                                                                                                                                                       0,
                                                                                                                                                                                                                                                                                                                                                                                                                         0
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CC probes used to screen for homozygous deletions. The region of deletion in CC cell line ASPC-1, and thereby to map deletions. The region of deletion in CC ASPC-1 was found to include a portion of the gene encoding a protein CC ASPC-1 was found to include a portion of the gene encoding a protein CC kinase, MKK4. MKK4 (also known as JNKK1 and SRK1) may be involved in a CC MAPK (mitogen-activated protein kinase) pathway for the signal CC transduction of cytokine-induced and stress-induced apoptosis. MKK4 is CC also involved in suppressing a variety of tumours. MKK4 is a dual CC also involved in suppressing a variety of tumours. MKK4 is a dual cc as MKK4 and p38 MAPKs are activated via dual phosphorylation on CC MAPKS. The JNK and p38 MAPKs are activated via dual phosphorylation on CC deletions often involve loss of a single allele, which is known as loss of heterozygosity (LOH), and the remaining allele is presumed to be non-CC deletions often involve loss of a single allele, which is known as loss CC deletions often involve loss of a single allele, which is known as loss CC deletions often involve loss of a single allele, which is known as loss CC deletions often involve loss of a single allele, which is known as loss CC deletions often involve loss of a single allele. Which is known as loss CC deletions often involve loss of a pre-axisting inherited mutation, or CC involve homozygous deletion of both alleles. LOH events commonly involve consorve because of a pre-axisting inherited mutation are cCC deletions spanning many megabases of DNA, while homozygous deletions are cCC deletions spanning many megabases of DNA, while homozygous deletions are cCC deletions spanning many megabases of DNA, while homozygous deletions are cCC deletions of the MKK4 locus or its expression product in a tissue sample cCC deletions of a predisposition to breast, pancreatic, colorectal and cCC deletion of the MKK4 locus or its expression product in a tissue sample cCC deletion for cancer to the deletion save been found in cell cell t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    breast cancer; pancreatic cancer; colorectal cancer; testicular cancer; drug screening; gene therapy; protein replacement therapy; mimetic; sequence tagged site; STS; probe; deletion; mapping; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           drug screening;
sequence tagged
protein may mutation in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Polynucleotides comprising all or a portion of the tumor suppressor gene MKK4 locus are useful for diagnosis, prognosis and therapy of human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Skolnick MH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13-JUN-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cancers
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                                     function for cancer therapy. MKK4 gene therapy, protein replacement therapy and protein mimetics that reconstitute the function of the protein may be used for therapy of human cancers which result from
         the MKK4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Perry WL,
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    gene
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                                                                                                      of the MKK4
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176 Н

TTTAGTAGAGATGGAGTTTCTC 197

Query Match Best Local : Matches

20;

Similarity

1.9%; 2 C; 7 G;

Score 18.8; Pred. No. 1

.3e+03

DB 1; Length 22,

0,

Mismatches

0,

Gaps

0

22

BP;

ψ Þ,

8 T; 0 U; 0 Other;

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RESULT 715
AAF29797/c
ID AAF297
XX AAF297
AC AAF297
XX O9-APR
XX Presen
XX Presen
XX Human;
KW Human;
KW presen
XX SHomo 8
XX Homo 8
                                                                                                                                                                                                                                                                                                                                                                  RESULT 714
AAF84349/c
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                                                                                                                                                                                                                                                                                                         Query Match
Best Local S
Matches 20
                                                                                                                                                                                                                                                                                                                                                                Sequence 22 BP; 5 A; 7 C; 3 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                      primer and a fluorescence-labelling probe. The method involves carrying out PCR on sample DNA, containing a drug-metabolising enzyme gene, together with PCR buffer, the normal forward primer, the reverse primer and the fluorescence-labelling probe (step A); and carrying out PCR on the sample DNA together with PCR buffer, the mutated forward primer, the reverse primer and the fluorescence-labelling probe (step B), and a step of comparing the result of step a with that of step b. The present sequence is a primer for human CYP2C18i, which was used to illustrate the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Detection of gene polymorphism of drug-metabolizing enzymes useful for diagnosis and testing comprises carrying out polymerase chain reaction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19-MAR-1999;
06-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 JP2001017185-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gene polymorphism;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human CYP2C18i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAF84349;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAF84349 standard; DNA; 22
                 Homo sapiens
                                            preseniline-1;
                                                          Human; PSEN1; Alzheimer's disease; polymorphism;
                                                                                                                   09-APR-2001
                                                                                                                                                                           AAF29797 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to a kit and method for the detection of gene polymorphisms of drug-metabolising enzyme genes. The kit contains a polymerase chain reaction (PCR) buffer solution containing DNA polymerase and NTP, a normal forward primer, a mutated forward primer, a reverse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 1; Page 13; 27pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2001-285409/30.
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                                                                                                                                               AAF29797;
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                                                                                                                                                                                                                                                  22
                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                             AGTGCTGGGATTACAGGCGTGA 882
                                                                                                                                                                                                                                                  AATGCTGGGATTACAGGCATGA 1
                                                                                                                                                                                                                                                                                                         Conservative
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                                                                                                                   (first
                                                                                      gene promoter PCR primer Prom22R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PCR
                                            chromosome 14;
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                                                                                                                                                                           DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     drug-metabolising enzyme; PCR primer; CYP2C18i;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   primer #5.
                                                                                                                 entry)
                                                                                                                                                                                                                                                                                                                     1.9%;
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                                                                                                                                                                           ВP
                                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                                      Score 18.8;
Pred. No. 1.
                                            PCR primer; 88
                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                        1.3e+03
                                                                                                                                                                                                                                                                                                                                     DB 1;
                                                                                                                                                                                                                                                                                                         2
                                                          diagnosis;
                                                                                                                                                                                                                                                                                                                                  Length 22;
                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        88
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RESULT 716
AAD31453/c
ID AAD31453
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S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             or susceptibility to Alzheimer's disease in humans, involving detecting genetic lesion in the presentiline-1 (PSENI) gene, found on chromosome if the genetic lesion is a polymorphism in the promoter or upstream regulatory region of the gene. The invention also describes transgenic animals which can be used to identify compounds useful in treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; Van Buchem's disease; genomic deletion; craniotubular hypertosis; autosomal recessive disorder; chromosome 17; chromosome 17q21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human chromosome 17 92Kb gene fragment amplifying PCR primer, WtlR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      31-MAY-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention describes a method for determining the presence or susceptibility to Alzheimer's disease in humans, involving detecting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 1; Page 45; 56pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            genetic lesion in the presenilin-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Determining wether a human subject has or is at risk of developing (early conset) Alzheimer's disease comprises detecting the presence/absence of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200079000-A1
                                                                           WPI; 2002-227089/28
                                                                                                                                                                                                                                                                  28-JUL-2000;
06-JUL-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200210455-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      bone dysplasia;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   174 TTTTTAGTAGAGATGGAGTTTC 195
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22
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                                                                                                                                   Ä
                                                                                                                                                                                   CELLTECH R
STRAEHLING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TTTTAGTAGAGACGGGGTTTC 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                     2000US-0221855P
2001US-0303386P
                                                                                                                                 Proll S,
                                                                                                                                                                                                                                                                                                                                                      2001WO-US023968.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         92Kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1.9%;
                                                                                                                                                                                         HAMPTON
                                                                                                                                                                                                               & D INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Van Broeckhoven
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      fragment; PCR primer; ss
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 18.8;
Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            gene.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 1;
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Methods for identifying subjects diseases associated with genomic

who are afflicted with or carriers of deletion(8), e.g. Van Buchem's diseas

Van Buchem's disease,

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FFXSX
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AAD31457/c
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to methods for distinguishing between individuals homozygous for and therefore afflicted with Van Buchem's disease, individuals heterozygous for and therefore carriers of Van Buchem's disease and individuals who are not afflicted with Van Buchem's disease comprise identifying a large genomic deletion in chromosome 17 at 17q21. The method is useful for identifying individuals who are afflicted with or carriers of diseases associated with one or more genomic deletion, particularly Van Buchem's disease, which is a rare autosomal recessive disorder that results in a bone dysplasia referred to a craniotubular hypertosis. The present sequence is a PCR primer used to amplify 92Kb gene fragment in human chromosome 17 at 17q21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           by determine chromosome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; Van Buchem's disease; genomic deletion; craniotubular hypertosis; autosomal recessive disorder; chromosome 17; chromosome 17q21; bone dysplasia; 92Kb gene fragment; PCR primer; 8s.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
The present invention relates to methods for distinguishing between individuals homozygous for and therefore afflicted with Van Buchem's disease, individuals heterozygous for and therefore carriers of Van Buchem's disease and individuals who are not afflicted with Van Buchems's disease comprise identifying a large genomic deletion in chromosome 17 at 17g21. The method is useful for identifying individuals who are afflicted
                                                                                                                              Methods for identifying subjects who are afflicted with or carriers of diseases associated with genomic deletion(s), e.g. Van Buchem's disease, by determining the presence of a deletion in the 92 kb region of human chromosome 17 at 17g21.
                                                                                                                                                                                                                                                                                                           28-JUL-2000;
06-JUL-2001;
                                                                                                                                                                                                                                                                                                                                                     30-JUL-2001; 2001WO-US023968
                                                                                                                                                                                                                                                                                                                                                                                    07-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                 WO200210455-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
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                                                                                                     Example
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                                                                                                                                                                                                                                                                              CELL-)
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                                                                                                                                                                                                                                                                CELLTECH R
STRAEHLING
                                                                                                     3; Page
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATCCTCCTGCCTCAGCCTCCCA 553
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTCTCTTGCCTCAGCCTCCCA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP; 7 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                           2000US-0221855P.
2001US-0303386P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                     Proll S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          g the presence
at 17q21.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              26; 109pp;
                                                                                                  26; 109pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17
                                                                                                                                                                                                                                                                HAMPTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 92Kb gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2 C; 10
                                                                                                                                                                                                                                   Paeper
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            of a deletion in the 92 kb
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Pred. No. 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    fragment amplifying
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670

TTGGCTCACTGCAACCTCTGCC

691

Query Match Best Local Matches

Similarity

1.9%;

Conservative

<u>,,</u>

Pred. No. 1.3 0; Mismatches Score 18.8;

No.

1.3e+03

DB 1;

Length

22;

Indels

0

Gaps

0

Sequence 22

BP;

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RESULT 718
ADL66998/c
ID ADL669
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                    The invention relates to a new composition which comprises at least one anti-DNA polymerases (anti-DNAP) antibody and/or at least one anti-DNAP polymerases (anti-DNAP) antibody, and at least one single strand binding protein (SSB) or at least two different SSBs. The compositions are useful for nucleic acid synthesis reactions or are generated during nucleic acid synthesis. The methods are useful for synthesis reactions. The methods are useful for synthesis to be used in amplifying nucleic acid molecules, in reverse transcription concleic acid molecules and in coupled or uncoupled reverse transcription/amplification. The present sequence is used in the exemplification of the present invention.
                                                                                                                                                                                                                         New compositions comprising one or more anti-reverse transcriptase antibodies, anti-DNA polymerases or single strand binding proteins, useful for synthesizing nucleic acids.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                      05-SEP-2002;
19-NOV-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA polymerase; anti-DNAP antibody; reverse transcriptase;
anti-RT antibody; single strand binding protein; SSB; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         03-JUN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADL66998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADL66998 standard;
                                                                                                                                                                                                                                                                                                                 Park K;
                                                                                                                                                                                                                                                                                                                                                                                                                  05-SEP-2003; 2003WO-US027705
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Multiplex PCR primer
                                                                                                                                                                                                 Example 4; Page 89; 201pp; English.
                                                                                                                                                                                                                                                                                     WPI; 2004-248479/23
                                                                                                                                                                                                                                                                                                                                          (INVI-)
                                                                                                                                                                                                                                                                                                                                                                                                                                            18-MAR-2004
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                                                                                                                                                                                                                                                                                                                                             INVITROGEN CORP
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                                                                                                                                                                                                                                                                                                                                                                      2002US-0408609P.
2002US-0427867P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
7 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA;
5 C;
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Pred. No.
2 T;
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  Other
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anscription of
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RESULT 719
RAA37708
ID AAA377
XX AAA377
XX AAA377
XX AAA118e
KW AAT18e
KW radiat
XX HOMO 8
PN WO2000
XX HOMO 8
PN WO2000
XX HOMO 8
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XX HOMO 9P 17-AUG
XX HOMO 9P 10-FEE
PR 06-DEC
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Best Local S
Matches 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  designated AS8 (also referred to as RSIAS8). The antisense inhibitors can be used in a method of the invention, for inhibiting cell proliferation. They can also be used in methods for inducing sensitivity to radiation and DNA damaging chemotherapeutics in an individual and in a method for prolonging survival in an individual with cancer. The methods and antisense molecules are useful for inhibiting cell proliferation, especially cancerous cell proliferation, for inducing sensitivity to radiation and DNA damaging chemotherapeutics in individuals and for prolonging survival in an individual with cancer. Kits for carrying out the methods may be used to diagnose and/or treat cancer and for addinctive there are there we are the services of the s
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Antisense radiation
Human; Rad51; antisense; drug screening; cancer; autoimmune disease;
                                                                          Human RAD51 antisense oligonucleotide, AS8.
                                                                                                                                                                 04-JUL-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Inhibiting cell proliferation useful for cancer therapy, comprises administering Rad51 inhibitor in vivo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAA37708 standard; DNA;
                                                                                                                                                                                                                                                                                                                              AAS01201 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 8; Page 26; 42pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22-NOV-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 06-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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                                                                                                                                                             (first entry)
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99US-00454495.
                                                                                                                                                                                                                                                                                                                          CDNA;
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Pred. No. 1.4e+03;
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Best Local S
Matches 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 such as cancer, autoimmune disease, arthritis, graft rejection, inflammatory bowel disease, proliferation induced after medical procedures such as surgery, angioplasty etc. in humans and animals
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The sequence represents the human Rad51 antisense oligonucleotide, AS8. The antisense oligonucleotide its used to study down-regulation of Rad5 protein in human brain, breast and prostate cells. Rad51 protein is defective in repair of damaged DNA, genetic recombination and the recombinational repair of DNA lesions, and plays a central role in cancer. Inhibiting cell proliferation involves administering to a cell
                                                                                                                                                                                                                                                                                        Tumour cell proliferation, RadS1 inhibitor; p53 protein; premature aging; hyperproliferative disorder; hodgkin's disease; squamous cell carcinoma; leukaemia; autoimmune disease; cancer; graft rejection; angioplasty; inflammatory bowel disease; immunosuppressive; gene therapy; arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAD43247;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAD43247 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rad51 antibody or its fragment. The Rad51 antibody or its fragment is useful for inhibiting cell proliferation, for treating disease states
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Inhibiting cell proliferation for treating arthritis, graft rejection, inflammatory bowel disease, cancer, proliferation induced after medical procedure, involves administering Rad51 antibody or its fragment to cell.
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06-DEC-1999;
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                                                                                                               modified_base
                                                                                                                                                                                                   Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                        Antisense oligonucleotide R51AS8.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14-NOV-2002 (first entry)
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20; Conserv
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                                                                                                                                                                                                                                                             phosphorothioate
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ilarity 90.9%;
Conservative
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99US-00455300.
                                                                                                                                          Location/Qualifiers
                                 note=
                                                                                 /*tag= a
                                                            base= OTHER
                           "Phosphorothioate backbone"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              <u>,</u>
                                                                                                                                                                                                                                                                backbone;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 18.8;
Pred. No. 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     for treating disease states
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Rad51
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Best Local S
Matches 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a method for inhibiting or reducing tumour cell proliferation in an individual in vivo. The method comprising contacting a tumour cell in vivo with a Rad51 inhibitor and a polynucleotide capable of expressing functional p53 protein. The method is useful for inhibiting or reducing tumour cell proliferation in an individual in vivo. The method is useful for treating hyperproliferative disorders, especially cancer (such as Hodgkin's disease, squamous cell carcinoma and leukaemia), premature aging, autoimmune disease, arthritis, graft rejection, inflammatory bowel disease, and proliferation induced after medical procedures such as surgery and angioplasty. The invention is useful in gene therapy. The present sequence is an antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              26-JAN-2001; 2001US-00771355
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Inhibiting/reducing tumor cell proliferation in individual in vivo, treating cancer, arthritis, involves contacting tumor cell in vivo Rad51 inhibitor, and polynucleotide expressing functional p53 prote
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Zarling DA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  26-JAN-2000; 2000US-0178561P
                                                                                                                                                                                            Human; heart failure-associated gene; hUNC93B1; ss; chromosome 11q13; SNP; single nucleotide polymorphism; heart failure; 12 transmembrane transporter; left ventricular diastolic heart failure; systolic heart failure; PCR; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 23 BP; 2 A; 13 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 5; 12pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        oligonucleotide used
 Andersson MK,
                                                                                 03-AUG-2001; 2001US-00922445
                                                                                                             04-MAR-2003.
                                                                                                                                        US6528268-B1
                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                  Human hUNC93B1 polymorphic region 24801/24941 primer #1
                                                                                                                                                                                                                                                                                                11-MAR-2004
                                                                                                                                                                                                                                                                                                                                                        ADH26585 standard;
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                                                      03-AUG-2001; 2001US-00922445
                          (SEQU-) SEQUENOM GEMINI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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                                                                                                                                                                                                                                                                                                (first entry)
 Berglund LGT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1.9%;
                                                                                                                                                                                                                                                                                                                                                         DNA;
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                            LTD.
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Pred No. 1.
Reneland
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 Adam GIR;
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                                                                                                                                                                                                               heart failure;
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Detection diagnosis

Page

13;

23pp;

Japanese

of human cytochrome p4501A2 gene poof metabolic activity polymorphism.

polymorphism

useful in

WPI; 1996-087678/09

AAT10907-T10910 are PCR primers used for the amplification of untranslated fragment of the the human cytochrome P4501A2 gene

gene

including

a

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RESULT 723
AAT10907/c
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Best Local S
Matches 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      diastolic heart failure in a human comprises detecting the presence or absence of an allelic variant at position 24941 of the heart failure associated gene appearing as ADH26544 (NUNC93B1, encoding a possible 12 transmembrane receptor protein) in the sample, where the presence or absence of the allelic variant is indicative of a predisposition to left ventricular diastolic heart failure in the human. The method further comprises determining the genotype of the human at position 24941 of ADH26544. The method is useful for diagnosing predisposition to left ventricular diastolic and systolic heart failure. The hUNC93B1 gene is located on chromosome 11q13. The present sequence is a PCR primer used to analyse the polymorphisms in the human heart failure associated gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             bisgnosing predisposition to left ventricular diastolic heart failure a human comprises detecting the presence or absence of an allelic variat position 24941 of hUNC93B1 gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI;
                                                                                                                                                                                                                                                                                                                                                                  Human cytochrome P4501A2 (CYPIA2) gene 5' UTR fragment PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to diagnosing predisposition to left ventricular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; SEQ ID NO 42; 53pp; English
                                                                                                                                                                                                                                06-JUL-1995;
                                                                                                                                                                                                                                                                                                         Synthetic.
                                                                                                                                                                                                                                                                                                                               Cytochrome P450; detection; diagnosis; polymorphism; substitution; metabolism; respiration; polymerase chain reaction; ss.
                                                                                                                                                                                                                                                                                                                                                                                            06-SEP-1996
                                                                                                                                                                                                                                                                                                                                                                                                                      AAT10907
                                                                                                                                                                                                                                                                                                                                                                                                                                              AAT10907 standard;
                                                                                                                                                                                                        06-JUL-1994;
                                                                                                                                                                                                                                                       18-JAN-1996.
                                                                                                                                                                                                                                                                                 WO9601328-A1
                                                                                                                      Fukui
                                                                                                                                          (SAKA ) OTSUKA PHARM CO LTD.
(KIMS/) KIM S.
(SHIN/) SHIN K.
(SHIN/) SHIN J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2004-088115/09
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                                                                                                                    Katsuragi
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           The invention relates to the isolation of polymorphic repeat sequences having the sequence (dC-dA)n.(dG-dT)n which can be used as genetic markers. Primers based on these sequences can be used to detect these repeats, especially for use in e.g paternity or maternity testing, human genetic analysis such as linkage analysis of genetic disease, commercial animal or plant breeding or pedigree analysis. Clones containing the repeat sequences were isolated by hybridisation of chromosome-specific phage libraries with a synthetic poly(dC-dA). (dG-dT) probe. Over 100 repeat blocks were isolated. The primers AAT65798-T66047 were used to PCF amplify the inserts from the isolated clones containing the repeat sequences. The primers AAT66010-1 were used to amplify the repeat sequence marker clone Mfd107 (AAT65778). (Updated on 25-MAR-2003 to
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                                                                                                                                                                                                                                                                                                                                                                                        Claim 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1997-042299/04.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Weber
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05-SEP-1991;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            hybridisation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Primer #1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25-MAR-2003
18-JUN-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             721 GCCTCCTGAGTAGCTGGGAC 740
                                                                                                                                                                                                                                                                                                                                                                                                                                               novel nucleic
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19; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                         of polymorphic genetic markers of the el nucleic acid mols. as primers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    to amplify repeat sequence marker Mfd107
field.)
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(first entry)
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91US-00754351.
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18-JUN-1997
                                                                                                                                       The invention relates to the isolation of polymorphic repeat sequences having the sequence (dC-dA)n. (dG-dT)n which can be used as genetic markers. Primers based on these sequences can be used to detect these repeats, especially for use in e.g paternity or maternity testing, huma genetic analysis such as linkage analysis. Clones containing the animal or plant breeding or pedigree enalysis. Clones containing the repeat sequences were isolated by hybridisation of chromosome-specific phage libraries with a synthetic poly(dC-dA). (dG-dT) probe. Over 100 repeat blocks were isolated. The primers AAT65798-T66047 were used to pamplify the inserts from the isolated clones containing the repeat sequences. The primers AAT66016-7 were used to amplify the repeat sequence marker clone Mfd110 (AAT65781). (Updated on 25-MAR-2003 to
                                                                                                    Sequence 20
                                                                                                                                                                                                                                                                                                                                        Detection of polymorphic genetic markers of the form (dC-dA)n(dG-dT)n using novel nucleic acid mols. as primers.
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05-SEP-1991;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       hybridisation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Polymorphism; repeat sequence; genetic marker; primer; amplification; PCR; polymerase chain reaction; paternity; maternity; human; pedigree; linkage analysis; genetic disease; animal; plant; breeding; locus;
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                                                                 Local
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ACCACAACACCTGGCTAATT 1
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                                                                                                                                field.)
                                                                                                     BP; 4 A;
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                                                                                                                                                                                                                                                                                                              13-14; 186pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     amplify repeat sequence marker Mfd110
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91US-00754351.
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                                                              1.9%;
95.0%;
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95.0%;
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Pred. No. 1.3e
0; Mismatches
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RESULT 727
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CC specifically a sense DPC4 sequence (particularly in the form of a vector, i.e. by gene therapy), but also an antisense sequence where DPC4 protein is over expressed or (b) mimics the activity of DPC4. DPC4 nucleic acid is also used as hybridisation probes for detecting presence/absence of thuman chromosome 18q21.1 fragments. When a homozygous deletion is CC detected in this region, an agent can be administered that accumulates within, or kills, only cells which contain such a deletion. This agent can be capablouring gene and lost by the deletion, i.e. it has a highly considered as hybridisation probes for other protein) encoded by a mention of the protein and a significant contains such a deletion.
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Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Deleted in Pancreatic Cancer locus 4 polypeptide - and related nucleic acids, used in diagnosis and treatment of proliferative diseases, e.g. cancer of pancreas or other organs.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DPC4; pancreatic cancer; deleted; locus 4; diagnosis; human;
tumour suppressor gene; proliferative disease; bile duct; bl
colorectal; cancer; Crohn's disease; colitis; PCR primer;
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19; Conser
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                                                                                                                                TCCCAAAGTGCTGGGATTAC 404
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                                                                                                                                                                                                                                 A;
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                                                                                                                                                                                              1.9%;
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                                                                                                                                                                                                                               G; 6 T; 0 U; 0 Other;
                                                                                                                                                                               Pred.
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Pred. No. 1.
                                                                                                                                                                 Mismatches
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                                                                                                                                                                                 1.3e+03
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AAV85762 standard; DNA;

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RESULT 728
AAV85840/c
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                                                                                                                                                                                                                                                                                        The present invention describes LRP5 (low density lipoprotein (LDL) receptor related protein, previously designated LRP-3). AAV85587 to AAV85822 represent exon primers used for obtaining LRP5 cDNA. Nucleic acid molecules (NAMS) encoding LRP5 can be used for determining if an individual is susceptible to insulin dependent diabetes mellitus (IDDM). The NAMS or proteins can be used for reducing triglyceride levels in the serum of an individual. Therapies that affect LRP5 may also be useful in the treatment of autoimmune diseases such as glomerulonephritis, diseases and disorders involving disruption of endocytosis and/or antigen presentation, cytokine clearance and/or inflammation, viral infection, pathogenic bacterial toxin contamination, elevation of free fatty acids or hypercholesterolemia, type 2 diabetes, osteoporosis, Alzheimer's disease and cardiovascular disease. Products from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LRP5; LDL-receptor related protein; LRP-3; IDDM; diagnosis; endocytosis; insulin dependent diabetes mellitus; autoimmune disease; glomerulonephritis; inflammation; viral infection; osteoporosis; hypercholesterolemia; Alzheimer's disease; low density lipoprotein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New isolated LDL-receptor related protein - used to develop products for treating, e.g. elevated triglyceride levels, diabetes, autoimmune disorders, inflammation or Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Todd JA, Hess JW, Caske:
Hey P, Kawaguchi Y, Mer
Phillips MS, Twells RCJ;
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                                         AAV85840;
                                                                                                                                                                                                                                                     Sequence 20
                                                                                                                                                                                                                                                                                 can also
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 12; Page 105; 200pp; English.
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             10-FEB-1999
                                                                    AAV85840
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         primer; ss.
                                                                                                                                                                 391 AGTGCTGGGATTACAGGCGT 410
                                                                                                                                       20
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                                                                                                                                                                                                          Similarity
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                                                                    standard; DNA; 20
                                                                                                                                                                                                                                                     BP; 5 A; 7 C; 3 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                               used for
                                                                                                                                                                                               Conservative
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             (first entry)
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97US-0048740P.
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Merriman
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                                                                                                                                                                                                          1.9%;
                                                                                                                                                                                                                                                                               detection, diagnosis and
                                                                     ΒP
                                                                                                                                                                                               0; Mismatches
                                                                                                                                                                                                              Pred.
                                                                                                                                                                                                                          Score 18.4;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cox
                                                                                                                                                                                                              No.
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Metzker ML,
                                                                                                                                                                                                             1.3e+03
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                                                                                                                                                                                                                             DB 1;
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                                                                                                                                                                                               Indels
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agawa Y;
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AAV85801
ID AAV
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AC AAV
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AC AAV
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LRP
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KW LRP
KW ins
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                                                                                                                                                        RESULT 729
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Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                     receptor related protein, previously designated LRP-3). AAV85823 to AAV85900 represent SNP primers used for obtaining LRP5 cDNA. Nucleic acid molecules (NAMs) encoding LRP5 can be used for determining if an individual is susceptible to insulin dependent diabetes mellitus (IDDM). The NAMs or proteins can be used for reducing triglyceride levels in the serum of an individual. Therapies that affect LRP5 may also be useful in the treatment of autoimmune diseases such as glomerulonephritis, diseases and disorders involving disruption of endocytosis and/or antigen presentation, cytokine clearance and/or inflammation, viral infection, pathogenic bacterial toxin contamination, elevation of free fatty acids or hypercholesterolemia, type 2 diabetes, osteoporosis, Alzheimer's disease and cardiovascular disease. Products from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LRP5; LDL-receptor related protein; LRP-3; IDDM; diagnos insulin dependent diabetes mellitus; autoimmune disease; glomerulonephritis; inflammation; viral infection; osteo
LRP5; LDL-receptor related insulin dependent diabetes
                                                                                                                                                                                                                                                                                                                Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New isolated LDL-receptor related protein - used treating, e.g. elevated triglyceride levels, dial disorders, inflammation or Alzheimer's disease.
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                                         LRP5 exon primer 58-10 lr.
                                                                      10-FEB-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 12; Page 110; 200pp; English.
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05-JUN-1997;
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                                                                                                                             AAV85801 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22-OCT-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          hypercholesterolemia; Alzheimer's disease; low density lipoprotein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention describes LRP5 (low density lipoprotein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       JA, Hess JW, Caske
P, Kawaguchi Y, Mer
lips MS, Twells RCJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sapiens.
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                                                                                                                                                                                                                   AGTGCTGGGATTACAGGCGT 410
                                                                                                                                                                                                  AGTGCTGGGATTACAGGCAT 1
                                                                                                                                                                                                                                                                                                              BP; 5 A; 7 C; 3 G; 5 T; 0 U; 0 Other;
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                                                                      (first
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97US-0048740P
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                                                                                                                                                                                                                                                                                                                                          for detection,
                                                                                                                                                                                                                                                                    1.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Caskey CT,
Merriman
                                                                     entry)
                                                                                                                              20
protein; LRP-3; IDDM; diagnosis;
mellitus; autoimmune disease;
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Pred.
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                                                                                                                                                                                                                                                                        18.4;
No. 1
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Metzker ML,
                                                                                                                                                                                                                                                                     1.3e+03
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                used to develop products for diabetes, autoimmune
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Nakagawa
                                                                                                                                                                                                                                                                                   Length
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                                                                                                                                                                                                                                                          Indels
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cagawa Y;
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              endocytosis;
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RESULT 730
AAV859
ID AAV859
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XX LRP5 S
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention describes LRP5 (low density lipoprotein (LDL) receptor related protein, previously designated LRP-3). AAV85897 to AAV85822 represent exon primers used for obtaining LRP5 cDNA. Nucleic acid molecules (NAMS) encoding LRP5 can be used for determining if an individual is susceptible to insulin dependent diabetes mellitus (LDDM). The NAMS or proteins can be used for reducing triglyceride levels in the serum of an individual. Therapise that affect LRP5 may also be useful in the treatment of autoimmune diseases such as glomerulonephritis, diseases and disorders involving disruption of endocytosis and/or antigen presentation, cytokine clearance and/or inflammation, viral infection, pathogenic bacterial toxin contamination, elevation of free fatty acids or hypercholesterolemia, type 2 diabetes, osteoporosis, Alzheimer's
               insulin dependent diabetes mellitus; autoimmune disease; glomerulonephritis; inflammation; viral infection; osteoporosis; hypercholesterolemia; Alzheimer's disease; low density liponrote. PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New isolated LDL-receptor related protein - used to develop products treating, e.g. elevated triglyceride levels, diabetes, autoimmune disorders, inflammation or Alzheimer's disease.
                                                                                                                                                                                                                AAV85879
                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
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05-JUN-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               glomerulonephritis; inflammation; viral infection; osteoporosis;
hypercholesterolemia; Alzheimer's disease; low density lipoprotein;
                                                                                LRP5; LDL-receptor related protein; LRP-3; IDDM; diagnosis; endocytosis;
                                                                                                                LRP5 SNP
                                                                                                                                                10-FEB-1999
                                                                                                                                                                                AAV85879;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            can also
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           disease and cardiovascular disease. Products from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1998-594573/50
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P. Kawaguchi
Twe]
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                                                                                                                primer 58-10
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                                                                                                                                                                                                                                                                                               GTTCACTGCAACCTCTGCCT
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                                                                                                                                                                                                                                                                                                                                                                                                                              20 BP; 3
                                                                                                                                                                                                                standard; DNA;
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97US-0048740P.
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Merriman
                                                                                                                                                                                                                                                                                                                                                                             1.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            detection,
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Pred. No. 1.
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RESULT 731
AAX90795/c
ID AAX907
XX AAX907
XX I 3-JAN
DT 13-JAN
DT Human
DX Human
XX PCR pp
KW Synthe
XX Synthe
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Matches 19; Conserv
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention describes LRP5 (low density lipoprotein (LDL) receptor related protein, previously designated LRP-3). AAV85923 to AAV85900 represent SNP primers used for obtaining LRP5 CDNA. Nucleic acid molecules (NAMs) encoding LRP5 can be used for determining if an individual is susceptible to insulin dependent diabetes mellitus (IDDM). The NAMs or proteins can be used for reducing triglyceride levels in the serum of an individual. Therapies that affect LRP5 may also be useful in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New isolated LDL-receptor related protein - used to detreating, e.g. elevated triglyceride levels, diabetes, disorders, inflammation or Alzheimer's disease.
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Hey P, Kawaguchi Y, Meri
Phillips MS, Twells RCJ;
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05-JUN-1997;
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                                                                                                                                                                                       13-JAN-2000
                                                                                                                                                                                                                                              AAX90795
                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
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                                                                                                                                                         Human 7SL RNA specific PCR primer-1.
                              WO9951255-A1
                                                         Homo sapiens
                                                                      Synthetic
                                                                                                                  synthesised;
                                                                                                                               PCR primer;
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MERCK & CO INC.
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                                                                                                                                                                                                                                              standard; DNA;
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                                                                                                                human 7SL RNA; amplify; human staufen cDNA; random hexamer primer; Superscript II rever
                                                                                                                                                                                                                                                                                                                                                                                                                                       BP; 3 A;
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97US-0048740P.
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95.0%;
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, Merriman
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                                                                                                                                                                                                                                              20 BP
                                                                                                                                                                                                                                                                                                                                                                              0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                           Score 18.4;
Pred. No. 1.
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Metzker ML, Nakagawa
                                                                                                                                                                                                                                                                                                                                                                                          1.3e+03
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                                                                                                                                                                                                                                                                                                                                                                                                          Length 20
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                                                                                                                  reverse tra
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cagawa Y;
                                                                                                                   transcriptase;
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RESULT 732
AAX86546
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Best Local S
Matches 19
PCR primers AAX86523-62 were used for amplification and sequencing of exons of the Rhesus D (RhD) antigen gene. The specification describes a RhD contributing to or indicative of the weak D phenotype, where the RhD polymucleotide carries at least one missense mutation as compared to the wild-type RhD, in its transmembrane and/or intracellular regions, especially in amino acid positions 2-16, 114-149, 179-225 or/and 267-397, with the proviso that the D antigen does not carry a single missense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present sequence is a PCR primer specific to human 7SL RNA. used to amplify human staufen (hStau) cDNA synthesised using ranhexamer primers and Superscript II reverse transcriptase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human staufen polypeptide useful inhibitors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Greider CW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           06-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 1; Page 25; 50pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1999-620168/53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                06-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                             04-OCT-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                  AAX86546;
                                                                                           Example; Page 33; 64pp; English.
                                                                                                                  Nucleic acid sequences correlated with Rhesus weak
for screening blood from donors and recipients for
                                                                                                                                                                                                                        23-JAN-1998;
                                                                                                                                                                                                                                              18-DEC-1998;
                                                                                                                                                                                                                                                                     29-JUL-1999.
                                                                                                                                                                                                                                                                                            WO9937763-A2
                                                                                                                                                                                                                                                                                                                   Homo
                                                                                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                                                                                                                                                                                                                                Allele;
                                                                                                                                                                                                                                                                                                                                                                                       Primer re617 used
                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAX86546 standard; DNA;
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                                                                                                                                                    WPI; 1999-469127/39
                                                                                                                                                                                                 (DRKB-) DRK BLUTSPENDEDIENST
                                                                                                                                                                                                                                                                                                                                                    primer;
                                                                                                                                                                                                                                                                                                                  sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      730 GTAGCTGGGACTACAGGCGC 749
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             l Similarity
19; Conserv
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                                                                                                                                                                                                                                                                                                                                                                Rhesus D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GTAGCTGGGACTACAGGCAC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP; 3
                                                                                                                                                                                                                                                                                                                                                       88
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                                                                                                                                                                          Wagner FF;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ļe
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                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
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                                                                                                                                                                                                                                              98WO-EP008319
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                                                                                                                                                                                                                                                                                                                                                               antigen;
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                                                                                                                                                                                                                                                                                                                                                                                      for amplification and sequencing of RhD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          20 BP
                                                                                                                                                                                                                                                                                                                                                                RhD;
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Pred. No. 1
                                                                                                                                                                                                   BADEN WUERTTEMBERG
                                                                                                                                                                                                                                                                                                                                                                  weak D
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                                                                                                                                                                                                                                                                                                                                                                phenotype; blood transfusion;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          .3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    for identifying telomerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length
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                                                                                                                    D phenotype, useful transfusion methods.
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Matches 19
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                                             hyperproliferative gene expression. The antisense compound is used to treat an animal having a disease or condition associated with mdm2, particularly a hyperproliferative condition, more particularly cancer, especially of the blood, brain, breast, lung or soft tissue, or
                                                                                                                                                                                        AAZ37473-Z37738 represent human mdm2 phosphorothioate oligonucleotides.
AAZ37471, AAZ37472, AAZ37739, AAZ37740 and AAZ37741 are used in the exemplification of the present invention. The present invention describes novel nucleotide antisense compounds, targetted to the 5' untranslated, translation termination codon, or 3' untranslated region of a nucleic acid encoding human mdm2, that modulates expression of human mdm2. The oligonucleotides mediate their effect by antisense inhibition of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Miraglia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human mdm2 gene; proliferation; tumour; phosphorothioate; p53; cancer; antisense; modulation; oligonucleotide; expression; inhibition; hyperproliferation; blood cancer; brain cancer; breast cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  07-JAN-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New antisense compounds used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1999-610754/52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       26-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Page 55;
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Matches 19; Conserv
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                                                                                                                                                                                                   AAZ37473-Z37738 represent human mdm2 phosphorothioate oligonucleotides. AAZ37471, AAZ37472, AAZ37739, AAZ37740 and AAZ37741 are used in the exemplification of the present invention. The present invention describes novel nucleotide antisense compounds, targetted to the 5' untranslated, translation termination codon, or 3' untranslated region of a nucleic acid encoding human mdm2, that modulates expression of human mdm2. The oligonucleotides mediate their effect by antisense inhibition of hyperproliferative gene expression. The antisense compound is used to treat an animal having a disease or condition associated with mdm2, particularly a hyperproliferative condition, more particularly cancer, especially of the blood, brain, breast, lung or soft tissue, or psoriasis, fibrosis, atherosclerosis or restenosis
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AAZ37716
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human mdm2 gene; proliferation; tumour; phosphorothicate; p53; cancer; antisense; modulation; oligonucleotide; expression; inhibition; hyperproliferation; blood cancer; brain cancer; breast cancer; ancer; breast cancer; brain cancer; senteroscierosis; lung cancer; soft tissue cancer; psoriasis; fibrosis; atherosclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         07-JAN-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (ISIS-) ISIS
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          TCTTGGCTCACTGCAACCTC 687
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                                                                                                          1.9%;
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                                                                                   Score 18.4;
Pred. No. 1.
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Pred. No. 1
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                                                                                                                                                            3 T;
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                                                                                                                                                            0 U;
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RESULT 735
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Best Local
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16-APR-1999;
15-JUL-1999;
                                                                                                                                                                                                                                                                                                                                   protein 5 isoform NUCPSSI encoding DNA sequence. UCP5 is involved in metabolism, and it may be involved in catalysing H+ leak, and therefore be involved in energetic inefficiency in vivo. The present invention relates to human and murine UCP5 nucleotide and protein sequences. There are three isoforms of human UCP5, hUCPSI, hUCPSS, hUCPSSI, and two isoforms of murine UCP5, mUCP5I and mUCP5S. The human UCP5 gene is located on chromosome 10q23-25. The nucleic acids encoding UCP5 can be used as hybridization probes, in chromosome and gene mapping, for the generation of antisense RNA and DNA and in the preparation of recombinant UCP5 proteins. UCP5 nucleic acids can be used in gene therapy for regulation of metabolic conditions. Upregulating or downregulating UCP5 activity in a mammal is used for modulating metabolic rate in the mammal, in particular upregulation of UCP5 activity stimulates an increase in markabolic rate in the mammal.
                                                                                                                                                                                        metabolic rate in an obese mammal. Other therapeutic applications associated with modulating UCP5 activity are treating symptoms associated with stroke, trauma (e.g. burn trauma), sepsis and infection. Detecting UCP5 activity can be used to assist predictions concerning metabolic conditions or risk for onset of obesity and as UCP5 may control the generation of reactive oxygen to diagnose impaired neural activity or neural degeneration. Anti-UCP5 antibodies can be used in diagnostic assays and for the affinity purification of UCP5 from recombinant cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAA28013 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Uncoupling protein 5; UCP5; metabolism; chromosome 10q23-25; H+ leak; metabolic rate; obesity; stroke; trauma; burn trauma; sepsis; infecti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Uncoupling protein isoform UCP5SI nucleotide sequence PCR primer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This sequence represents a PCR primer specific for the human uncoupling protein 5 isoform hUCP5SI encoding DNA sequence. UCP5 is involved in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Adams S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              03-NOV-1999;
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                                                                                                                                          Sequence 20 BP; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Isolated nucleic acid encodes human uncoupling protein 5 useful in diagnostic assays and treatment of obesity, stroke, trauma, sepsis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (GETH ) GENENTECH INC
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                    866 TGGGATTACAGGCGTGAGCC 885
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                                                                         19;
                                                                                      Similarity
                                                                                                                                                                              or natural sources
 TGGGATTACAGGCATGAGCC 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pan J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Page 37; 90pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
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99US-0129583P.
99US-0143886P.
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                                                                                      1.9%;
                                                                                                                                          4 C; 7 G; 4 T; 0 U; 0 Other,
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                                                                       0;
                                                                                        Score 18.4; DB 1;
Pred. No. 1.3e+03;
                                                                         Mismatches
                                                                                                       Length 20;
                                                                         Indels
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RESULT 737 AAZ52253

AAZ52253

AAZ52253 standard; DNA; 20

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AAALI 943
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                                                                                                                                                                                                                                                                                                                                                                       CC in length, targeted to a nucleic acid encoding a human MDMX.

CC specifically hybridizes with and inhibits the expression of human MDMX.

CC specifically hybridizes with and inhibits the expression of human MDMX.

CC file products of the invention have anticarcinogen, antiinflammatory and

CC antiinfectious activity. Synthesized chimeric oligonucleotides targeted

CC to human MDMX, 20 nucleotides in length, composed of a central gap region

CC consisting of ten 2'-deoxynucleotides flanked on both sides by 5-

CC nucleotide wings were tested for antisense inhibition of MDMX expression.

CC Results of real-time quantitative polymerase chain reaction (PCR) showed

CC 10 out of the 159, 20 base pair sequences, all fully defined in the

CC specification, demonstrated at least 30% inhibition of MDMX expression.

CC modulation, particularly inhibition of MDMX expression.

CC modulation, particularly inhibition of MDMX expression, and may be used

CC disease or condition associated with expression of MDMX. The antisense

CC oligonucleotides may also be used as research reagents or kits, and as

CC distinguish between functions of various members of a biological pathway,

CC and as prophylaxis, e.g. to prevent or delay infection, inflammation or

CC tumor formation. AAA11781-AL1945 represent antisense oligonucleotides
                                                                                                                                       Matches
                                                                                                                                                                 Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New antisense oligonucleotides targeting nucleic acids encoding human MDMX useful for inhibiting MDMX expression and for treating diseases associated with MDMX expression e.g. tumor formation, inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAA11943 standard; DNA; 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human MDMX antisense oligonucleotide #31223
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                                                                                                                                                                                                                                                                               Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This invention describes a novel antisense compound (I),
                                                                                                                                                                                                                                                                                                                                                 described in the method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (ISIS-) ISIS PHARM INC.
                                               648 GCTGGAGTGCAGTGGCGCAA 667
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                                                                                                                                                                                                                                                                           BP; 4 A; 4 C; 8 G; 4 T; 0 U; 0 Other;
                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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                                                                                                                                           0; Mismatches
                                                                                                                                                                            Score 18.4; DB 1
Pred. No. 1.3e+03
                                                                                                                                                                                                                                                                                                                                                     invention
                                                                                                                                                                                                                 DB 1;
                                                                                                                                                                                                             Length 20;
                                                                                                                                               Indels
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RESULT 738
AAF31822/c
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AXU
                                                                                                                                                                                                                                                                                                                                       The present sequence is a primer 2C12502 used for sequencing a cDNA CC corresponding to an expressed sequence tag identified in a human lung CC library, to obtain full length clone of polynucleotide encoding stomach CC protein zsig28. The zsig28 gene is located at 3g22.1-3g22.2 region of CC human chromosome 3. The zsig28 protein shows homology to a diverse family CC murine oligodendrocyte-specific protein (OSD) and rat androgen-withdrawal CC apoptosis protein RVP.1. It is thought to be a cell-cell signalling CC molecule, a growth factor receptor or extracellular matrix associated CC protein with growth factor receptor or extracellular matrix associated CC apoptotic cellular pathway. The protein may act as an anti-microbial CC agent and may bind toxins produced by bacteria which cause food CC agonists are useful for promoting apoptosis in cells over-expressing CC asig28 e.g. in cancer cells. They are also useful for stimulating cell CC growth or differentiation. Altered levels of zsig28 protein in a test CC sample such as saliva, serum, sweat or biopsy can be monitored as an CC indication of digestive function, gastric ulcer or cancer. zsig28 CC expression can be used as a differentiation marker to determine the stage of tumour or cell maturity, particularly in epithelial cells. CC polynucleotides encoding zsig28 can be used in gene therapy applications cc. in the protein and cancer in the test contracts of the protein service.
                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                              Query Match
Best Local Similarity
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AAF31822;
                                    AAF31822 standard; DNA; 20
                                                                                                                                                                                                                                                                                           Sequence 20 BP; 5 A; 3 C; 8 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2000-271379/23.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            14-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  23-MAR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200015659-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                diagnosis; prevention; treatment; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            18-JUL-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 1; Page 121; 121pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (ZYMO ) ZYMOGENETICS INC
                                                                                                                                                                                                                                                                                                                               increase or inhibit zsig28 activity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       isolated polynucleotide encoding a stomach zsig28 polypeptide used diagnosis, prevention and treatment of stomach disorders caused by
                                                                                                                                                                          864
                                                                                                                                                                                                              19;
                                                                                                                                                                        GCTGGGATTACAGGCGTGAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           В
                                                                                                                                   GCTAGGATTACAGGCGTGAG 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       gastric ulcers or cancer.
                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Foley KP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       98US-00154444
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            99WO-US021023
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   for sequencing human stomach protein zsig28
                                                                                                                                                                                                                              1.9%;
                                    ВР
                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                Score 18.4; DB 1;
Pred. No. 1.3e+03;
                                                                                                                                                                        883
                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                  DB 1; Length 20;
                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CDNA
                                                                                                                                                                                                            0,
                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                            0
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ARESULT 739
ARF31823/C
ID AAF518
XX AAF618
XX IO-APR
AC AAF618
XX IO-APR
XX 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present sequence is one of a number of antisense compounds of 8 to 30 nucleobases in length that have been designed to target a 5'untranslated region, start codon, coding region of antiranslated region of the human receptor activator of NF-kappaB (RANK). The antisense compounds specifically hybridise with and inhibit the expression of RANK. The antisense oligonucleotides are useful for inhibiting the expression of human RANK in human cells or tissues. They can be utilised for diagnostics, therapeutics for the treatment of diseases associated with the expression of RANK, prophylaxis e.g. to prevent or delay infection, inflammation or tumour formation, and as research reagent. The antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel antisense compounds capable of modulating expression of human receptor activator of NF-kappaB useful for diagnosis, prophylaxis and treatment of diseases associated with expression of RANK.
                                                                                                                                                                                                                                   Human; cytostatic; receptor activator
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; cytostatic; antiinflammatory; antisense oligonucleotide; cancer; receptor activator of NF-kappaB; RANK; infection; inflammation; ss.
   05-NOV-1999;
                                                         09-JAN-2001.
                                                                                                                   US6171860-B1
                                                                                                                                                                                                                                                                                                                     Human RANK antisense oligonucleotide, SEQ ID NO: 81.
                                                                                                                                                                                                                                                                                                                                                                                  10-APR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                              AAF31823;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAF31823 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           compounds are safely and effectively administered to humans
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 14; Col 44; 40pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2001-136876/14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Baker BF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         05-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               05-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      09-JAN-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US6171860-B1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10-APR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             843
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CCTGCCTCGGCCTCCCAAAG 862
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CCAGCCTCGGCCTCCCAAAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20 BP; 2 A; 4 C; 10 G; 4 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Cowsert LM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     antisense oligonucleotide,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         99US-00435296
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  99US-00435296.
   99US-00435296.
                                                                                                                                                                                                                                   antiinflammatory; antisense oligonucleotide;
of NF-kappaB; RANK; infection; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 18.4; DB 1;
Pred. No. 1.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ـــ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        80
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                    cancer;
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RESULT 740
AAD14819/c
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                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 5 A; 8 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                               region, start coding region or 3'untranslated region of the human receptor activator of NP-kappaB (RANK). The antisense compounds specifically hybridise with and inhibit the expression of RANK. The antisense oligonucleotides are useful for inhibiting the expression of human RANK in human cells or tissues. They can be utilised for diagnostics, therapeutics for the treatment of diseases associated with the expression of RANK, prophylaxis e.g. to prevent or delay infection, inflammation or tumour formation, and as research reagent. The antisense compounds are safely and effectively administered to humans
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel antisense compounds capable of modulating expression of human receptor activator of NF-kappaB useful for diagnosis, prophylaxis and treatment of diseases associated with expression of RANK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Baker BF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 05-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 14; Col 44; 40pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present sequence is one of a number of antisense compounds of 8 to 30 nucleobases in length that have been designed to target a 5'untranslated
                                                                                                                                                           Homo sapiens
Synthetic.
                                                                                                                                                                                                       antisense therapy; diabetes; hyperproliferative disorder; inflamma neurological disorder; tumour; haematopoietic disorder; infection;
                                                                                                                                                                                                                                             Human glycogen synthase kinase 3 alpha antisense oligo ISIS #116660
                                                                                                                                                                                                                                                                  01-NOV-2001
                                                                                                                                                                                                                                                                                                       AAD14819
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (ISIS-) ISIS PHARM INC
modified_base
                                                     modified_base
                                                                                           modified_base
                                                                                                                              modified_base
                                                                                                                                                                                      phosphorothioate
                                                                                                                                                                                                 hyperproliferative disorder;
                                                                                                                                                                                                                           Human; glycogen synthase kinase 3 alpha; antidiabetic;
                           modified_base
                                                                                                                                                                                                                                                                                                                                                                392 GTGCTGGGATTACAGGCGTG 411
                                                                                                                                                                                                                                                                                                                                                     20
                                                                                                                                                                                                                                                                                                                                                                                          19;
                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                        standard;
                                                                                                                                                                                                                                                                                                                                                     GTACTGGGATTACAGGCGTG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cowsert LM;
                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  99US-00435296
                                 /*tag=
/mod_b;
                                                                                                             /*tag= a
/mod_base=
                /*tag=
                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                        backbone;
       mod_base=
                                                                                                     note=
                                                                note=
                                                                                                                                                                                                                                                                                                        DNA;
                                                                                                                                                                                                                                                                                                                                                                                                   1.9%;
                                     base=
                                                                 "Methoxyethyl
                                                                                                     "Phosphorothioate
                                                                                                                                                                                                                                                                                                        20
                  Œ
                                     m5c
                                                                                                                                                                                         88.
                                                                                                                                                                                                                                                                                                        ВP
                                                                           OTHER
                                                                                                               OTHER
                                                                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                developmental disorder; antisense;
                                                                                                                                                                                                                                                                                                                                                                                                     Score 18.4; DB 1;
Pred. No. 1.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                  residues"
                                                                                                      backbone"
                                                                                                                                                                                                                                                                                                                                                                                                              DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                              Length
                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                            cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                              20;
                                                                                                                                                                                                                   inflammation;
                                                                                                                                                                                                                                                                                                                                                                                           0,
                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                           0
RESULT 741
AAD14817/c
ID AAD148
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Query Match
Best Local S
Matches 19
                                                                                                                                                                                      length targetted to a nucleic acid encoding glycogen synthase kinase 3 alpha. The antisense compound specifically hybridises with and inhibits the expression of glycogen synthase kinase 3 alpha. The antisense compound is useful for the treatment of a diseases associated with glycogen synthase kinase 3 alpha such as diabetes, a neurological disorder, a haematopoietic disorder, a hyperproliferative disorder or a developmental disorder. The antisense compounds may also be used prophylactically to prevent or delay infection, inflammation or tumour formation. The present sequence is a phosphorothicate antisense oligonucleotide targetted to human glycogen synthase kinase 3 alpha
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Antisense compound 8 to 30 nucleobases in length comprising a that is targeted to a nucleic acid molecule encoding glycogen kinase 3 alpha, useful for the treatment of e.g. diabetes and hyperproliferative disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     modified_base
                                                                                                                                       Sequence 20
                                                                                                                                                                            genomic
                                                                                                                                                                                                                                                                                                                                                                              The invention relates to an antisense compound 8 to 30 nucleobases in
                                                                                                                                                                                                                                                                                                                                                                                                               Example 15; Page 84; 115pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2001-442247/47.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (ISIS-) ISIS PHARM INC.
                                   863
20
                                                                       19;
                                                                                                                                                                              DNA
                                                                                      Similarity
                        TGCTGGGATTACAGGCGTGA 882
 TGCTGGGATTACAGGGGTGA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mckay R,
                                                                                                                                          B₽;
                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /*tag= t
/mod_base=
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/mod_base=
19
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pom_
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/mod_base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 *tag=
                                                                                                                                         ð
                                                                                  1.9%;
                                                                                                                                          9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                base=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Butler MM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     base=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "Methoxyethyl residues"
                                                                                                                                          C; 2G; 4T; 0U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           m5c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                m5c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       m5c
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                                                                     0,
                                                                                        Score 18.4; DB 1
Pred. No. 1.3e+03
                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Wyatt JR;
                                                                                                                                            0 Other;
                                                                                                         DB 1;
                                                                                                         Length
                                                                           Indels
                                                                                                           20;
                                                                       0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       compound
synthase
                                                                       Gaps
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AAD14817 standard; DNA; 20

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The invention relates to an antisense compound 8 to 30 nucleobases in length targetted to a nucleic acid encoding glycogen synthase kinase 3 alpha. The antisense compound specifically hybridises with and inhibits the expression of glycogen synthase kinase 3 alpha. The antisense compound is useful for the treatment of a diseases associated with glycogen synthase kinase 3 alpha such as diabetes, a neurological disorder, a haematopoietic disorder, a hyperproliferative disorder or a developmental disorder. The antisense compounds may also be used prophylactically to prevent or delay infection, inflammation or tumour
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       antisense therapy; diabetes; hyperproliferative disorder; inflamma neurological disorder; tumour; haematopoietic disorder; infection; hyperproliferative disorder; developmental disorder; antisense;
                                                                                                                                                          Antisense compound 8 to 30 nucleobases in length comprising a that is targeted to a nucleic acid molecule encoding glycogen kinase 3 alpha, useful for the treatment of e.g. diabetes and
                                                                                                                                                                                                         WPI; 2001-442247/47.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human glycogen
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                                                                                                                        Example 15; Page 83; 115pp; English.
                                                                                                                                               hyperproliferative disorders.
                                                                                                                                                                                                                                                                             21-JAN-2000; 2000US-00488856
                                                                                                                                                                                                                                                                                                    16-JAN-2001; 2001WO-US001411
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ID AAK951
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11-JAN-2000;
02-MAY-2000;
                                                                                                                                                                                                                                                                                                     The invention relates to primers for synthesising full length cDNA clones. 830 cDNA molecules encoding a human protein have been isolated and nucleotide sequences of 5'- and 3'-ends of the cDNA molecules have been determined. Primers for synthesising the full length cDNA are useful for clarifying the function of the protein encoded by the cDNA. The full length clones were obtained by construction of full length enriched cDNA length clones were obtained by construction of full length enriched cDNA interface that were contacted by the cDNA construction of the protein encoded by the cDNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           genomic DNA
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                                                                                                                                                                               Sequence
                                                                                                                                                                                                                              clone
                                                                                                                                                                                                                                                                         enable the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                830 Primers useful for synthesizing full length cDNA clones and their use
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ota T,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Wakamatsu
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (-ITHH)
                                                                                                                                                                                                                                          libraries that were synthesised by the oligo-capping method. The primers enable the production of the full length cDNA easily without any special methods. The present sequence is a primer used to amplify a human cDNA
                                                                                                               Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          genetic
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19; Conserv
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                                          CCTGAGTAGCTGGGACTACA 744
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2000JP-00118774.
2000JP-00183765.
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                                                                                                                                                                                                                              the invention
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95.0%;
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Pred. No. 1
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                                                                                       Mismatches
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K, Kojima
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1 S, Otsuki
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T, Ko
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10-JUL-2000;
10-JUL-2000;
19-SEP-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention provides a database of human samples obtained from healthy individuals which can be used to identify polymorphic genetic markers. Data obtained for the database can be used to sort the samples by parameters such as age, sex and ethnicity. This is useful in linking markers with diseases, susceptibility to infection and drug responses. The present primer was used in an assay to demonstrate the uses of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         producing a database for identifying polymorphic genetic markers, comprises obtaining data relating to members of a healthy populat entering the information into a database.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human AKAP10 coding
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                                                                                                                                                                                            17-MAY-2001
                                                                                                                                                                                                                                    AAF92892
                                                                                                                                                                                                                                                                           AAF92892 standard;
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onse; PCR primer; ss.
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; 2000US-0217658P.
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RESULT 745
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15-MAR-2000;
23-JUN-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention relates to a method for treating a patient diagnosed as having a lower than normal high density lipoprotein-cholesterol (HDL-C) level, a higher than normal triglyceride level, or a cardiovascular disease, involving administering a compound that modulates LXR- or RXR-mediated transcriptional activity or ABC1 expression or activity. The LXR gene product may be used in an assay to identify compounds useful for the treatment of a disease or condition selected a lower than normal HDL cholesterol level, a higher than normal LDL cholesterol level, a higher than normal triglyceride level, and a cardiovascular disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Treating a lower than normal high density lipoprotein-cholesterol (HDL-C) level, a higher than normal triglyceride level, or a cardiovascular disease, by administering a compound that modulates LXR- or RXR-mediated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Fig 3; 317pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hayden MR,
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                                                                                                                                                                                                                                                                                                                                                                    Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                         Cancer associated protein; FOR gene; FRA16D; fragile site; aphidicolin; chromosomal rearrangement; cancer; splice variant; DNA instability;
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                       Richards R,
Nancarrow J,
                                                                                                                                                                                                                  15-DEC-2000; 2000WO-AU001539
                                                                                                                                                                                                                                                                  21-JUN-2001.
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                                                                                                                                            16-DEC-1999;
19-APR-2000;
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oxidoreductase; neoplasia; l
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2000US-0213958P.
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2000AU-00007025.
                         Ried K, F
Woollatt
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                                                                                                                                                                                                                                                                                                                                                                                                                       splice variant;
PCR primer; ss.
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                                                   Mangelsdorf M,
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Nancarrow

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Baker

WPI; 2001-398151/42

PCR primers AAH27888-AAH28055 represent PCR primers used to amplify identify minimal deletions in the human FRA16D oxidoreductase (FOR)

and gene

Example 1; Page 46; 150pp; English.

Novel isolated 16q23.2 nucleic acid molecule, FRA16D oxidoreductase gene associated with FRA16D site, useful for early diagnosis and assessment of risk of cancers associated with the FRA16D region.

(FOR)

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RESULT 746
AAF80892/c
ID AAF808
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XX Human
XX Antise
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Matches 19
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     The present invention relates to an antisense compound 8-30 nucleobases in length targeted to nucleobases 1-308 of the 5' untranslated region, 1776-1806 of the translation termination codon region or 1818-2370 of the 3' untranslated region of a nucleic acid molecule encoding human mdm-2.
                                                                                                                                                                                                  Novel antisense compound 8-30 nucleobases in 10 acid molecule encoding human mdm-2 useful for mof human mdm-2 and reducing hyperproliferation
                                                                                                                                                                                                                                                                                                                                                                               Miraglia LJ,
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                                                                                                                                            77pp;
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                                                                                                                                                                                                                                                                                                                                                                               Graham
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 18.4;
Pred. No. 1.
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                                                                                                                                                                                                                                                                                                                                                                               Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               .3e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1;
                                                                                                                                                                                                        in length targeted to a nucleic
for modulating the expression
ation of human cells.
                                                                                                                                                                                                                                                                                                                                                                               Cowsert LM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0
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RESULT 748 AAH38602

AXU

AAH38602

DNA;

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AAH38602;

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RESULT 747
AAF80870/c
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                      S
                                                Query Match
Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local S
Matches 19
                                                                                                                                   The present invention relates to an antisense compound 8-30 nucleobases in length targeted to nucleobases 1-308 of the 5' untranslated region, 1776-1806 of the translation terminon codon region or 1818-2370 of the 3' untranslated region of a nucleoic acid molecule encoding human mdm-2. The invention is useful for reducing hyperproliferation of human cells, modulating the expression of mdm2 in human cells or tissues or in vitro.
                                                                                                                                                                                                                                              Novel antisense compound 8-30 nucleobases in length targeted to a nucleic acid molecule encoding human mdm-2 useful for modulating the expression of human mdm-2 and reducing hyperproliferation of human cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention is useful for reducing hyperproliferation of human cells, modulating the expression of mdm2 in human cells or tissues or in vitro. The hyperproliferative disorder includes cancer or psoriasis
                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       02-MAY-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAF80870;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAF80870
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                          The
                                                                                                                                                                                                                        Example 9; Col 31; 77pp; English.
                                                                                                                                                                                                                                                                                                    WPI; 2001-190948/19
                                                                                                                                                                                                                                                                                                                           Miraglia
                                                                                                                                                                                                                                                                                                                                                                            26-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                     26-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     US6184212-B1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Antisense; mdm2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human mdm2
                                                                                                                                                                                                                                                                                                                                                     (ISIS-) ISIS
                                                             Local Similarity
                                                                                                                        hyperproliferative disorder includes cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           868
                        899
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20 GGATTACAGGCATGAGCCAC 1
                                                 19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                           ٢
                TCTTGGCTCACTGCAACCTC 687
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GGATTACAGGCGTGAGCCAC 887
                                                                                                 20 BP; 6 A; 4 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20 BP; 3 A; 6 C; 5 G; 6 T; 0 U; 0 Other;
 TCTTGGCTCACTGCAAGCTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             phosphorothioate oligonucleotide
                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                    PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                           Nero P,
                                                                                                                                                                                                                                                                                                                                                                              98US-00048810
                                                                                                                                                                                                                                                                                                                                                                                                     9908-00280805
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    hyperproliferation;
                                                           1.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1.9%;
                                                                                                                                                                                                                                                                                                                            Graham MJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20
                                                <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 18.4; DB 1;
Pred. No. 1.3e+03;
                                                             Score 18.4;
Pred. No. 1.
 _
                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                              Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cancer; psoriasis;
                                                           1.3e+03
                                                                           DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        or tissues or in vitro.
                                                                        Length
                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                          psoriasis
                                                                                                                                                                                                                                                                                                                              Z
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                                                                           20
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                                                 Gaps
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                                                 0;
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Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis; inflammation; forensic investigation; paternity analysis; PCR primer; ss.

SNP specific lower PCR primer

SEQ ID

1398.

14-AUG-2001

(first entry)

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RESULT 749
AAH37610/c
ID AAH376
XX
                                                                                                                                                                                                                                                                                                                                        CC primer extension (SNPE) primers, and the sequences of regions flanking concludes kits for determining the presence or absence of a SNP, using the colligonucleotides of the invention. The PCR primers are used to amplify a colligonucleotides of the invention. The PCR primers are used to amplify a colligonucleotides are useful for genotyping a nucleic acid sample by performing a single-nucleotide primer extension reaction. The CC oligonucleotides are useful for determining the presence, absence or colligonucleotides are useful for determining the presence, absence or colligonucleotides are useful for determining the presence, absence or colligonucleotides are useful for determining the presence, absence or colligonucleotides are useful for determining the presence, absence or colligonucleotides are useful for determining the presence, absence or colligonucleotides are useful for determining the presence, absence or colligonucleotides are useful for determining the presence, absence or colligonucleotides are useful for determining the presence of colligonucleotides are useful for determining the presence of colligonucleotides are useful for determining the presence or colligonucleotides are useful phenotypic trait suspected of being cased by one or more SNPs. Phenotypic traits include diseases e.g. colligonucleotides and acute intermittent porphyria. Phenotypic colligonucleotides are susceptibility to multifactorial colligonucleotides are susceptibility to multif
                                                                                                                           밁
                                                                                                                                                                                                      Query Match
Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nuc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             13-OCT-2000; 2000WO-US028436
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; Page 57; 83pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             acid sample.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Picoult-Newburg L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2001-290930/30
                                                                                                                                                                  869
                                                                                                                                                                                                      l Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORCHID BIOSCIENCES INC.
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                                                                                                                                                                GATTACAGGCGTGAGCCACC 888
                                                                                                                           GATTACAGGCATGAGCCACC 20
                                                                                                                                                                                                                                                                                         BP;
                                                                                                                                                                                                                                                                                                                                SNP containing
                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      99US-0160096P.
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                                                                                                                                                                                                                          1.9%;
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C;
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                                                                                                                                                                                                                                                                                                                                DNA sequence
                                                                                                                                                                                                         0,
                                                                                                                                                                                                                                                                                      G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                             Score
                                                                                                                                                                                                                            Pred.
                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                            18.4;
No. 1
                                                                                                                                                                                                                            3e+03
                                                                                                                                                                                                                                                DB 1;
                                                                                                                                                                                                                                                Length 20
                                                                                                                                                                                                           Indels
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AAH37610 standard; DNA;

20

ΒP

RESULT 750 AAH40090 ID AAH400

AAH40090 standard; DNA; 20

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188

GGAGTTTCTCCATGTTGGTC 207 GGGGTTTCTCCATGTTGGTC 1

0

Query Match Best Local S Matches 19

Similarity

1.9%;

Score 18.4; Pred. No. 1 Mismatches

1.3e+03;

DB 1;

Length Indels

0

Gaps

0

Conservative

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Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide CC primer extension (SNPE) primers, and the sequences of regions flanking CC sites of single nucleotide polymorphisms SNPs. The present invention CC includes kits for determining the presence or absence of a SNP, using the CC oligonucleotides of the invention. The PCR primers are used to amplify a CC performing a single-nucleotide primer is used as a genotyping primer. CC The oligonucleotides are useful for genotyping a nucleic acid sample by CC performing a single-nucleotide primer extension reaction. The CC individuals, having a pathological phenotypic trait suspected of being CC assess by association analysis the genotype of an individual or group of cc individuals, having a pathological phenotypic trait suspected of being CC agammaglobulinaemia, diabetes insipidus, Lesch-Nyhan syndrome, muscular CC dystrophy, familial hypercholesterolaemia, polycystic kidney disease, cc craits also include symptoms of or susceptibility to multifactorial CC disease of which a component is or may be genetic such as autoimmune cd iseases, including, rheumatoid arthritis, multiple sclerosis, cancer, nervous system diseases and infection by pathogenic CC microorganism. The method is also useful in forensic investigations and ccc for a human SNP containing DNA sequence represents a PCR primer specific ccc for a human SNP containing DNA sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPB; genotyping; agammaglobulinaemia; diabettes instpidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis; inflammation; forensic investigation; paternity analysis; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; Page 52; 83pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13-OCT-2000; 2000WO-US028436
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         26-APR-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAH37610;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Picoult-Newburg L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15-OCT-1999;
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Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (ORCH-) ORCHID BIOSCIENCES INC
BP;
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  7 C; 4 G; 1 T; 0
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  ū,
    0
    Other,
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AAH40090;

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                                                                                                                                                                                                                                                                                     Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide CC primer extension (SNPB) primers, and the sequences of regions flanking CC sites of single nucleotide polymorphisms SNPs. The present invention CC includes kits for determining the presence or absence of a SNP, using the CC includes kits for determining the presence or absence of a SNP, using the CC SNP flanking sequence, the SNPE primer is used as a genotyping primer. CC The oligonucleotides are useful for genotyping a nucleic acid sample by CC performing a single-nucleotide primer extension reaction. The CC oligonucleotides are useful for determining the presence, absence or CC identity of a SNP and for genotyping nucleic acid samples for eg. to assess by association analysis the genotype of an individual or group of CC individuals, having a pathological phenotypic trait suspected of being CC auseed by one or more SNPs. Phenotypic traits include diseases e.g. CC agammaglobulinaemia, diabetes insipidus, Lesch-Nyhan syndrome, muscular CC osteogenesis imperfecta and acute intermittent porphyria. Phenotypic traits also include symptoms of or susceptibility to multifactorial CC disease of which a component is or may be genetic such as autoimmune inflortence of a component is or may be genetic such as autoimmune conficer nervous system diseases and infortion by nathogenic conficer nervous system diseases and infortion by nathogenic conficer nervous system diseases and infortion by nathogenic
                                                                               Query Match
Best Local Similarity
Matches 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; polycystic kidney disease; rheumatoid arthritis; multiple sclerosis; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis;
                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; Page 64; 83pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                13-OCT-2000; 2000WO-US028436
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SNP specific lower PCR primer SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14-AUG-2001
                                                                                                                                                                                                                                                    inflammation, cancer, nervous system diseases and infection by pathogen microorganism. The method is also useful in forensic investigations and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Picoult-Newburg L,
                                                                                                                                                                                                                                paternity analysis. The present sequence represents a PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (ORCH-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2001-290930/30
                                       870
1 ATTACAGGCGTGAGCCACAA 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORCHID
                                                                                                                                                                  20 BP; 7
                                       ATTACAGGCGTGAGCCACCA 889
                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
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                                                                                                                                                                                                         containing
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                                                                                                   1.9%;
                                                                                                                                                                  5 C; 5 G; 3 T; 0 U; 0 Other,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      investigation; paternity analysis; PCR primer; ss
                                                                                                                                                                                                            DNA sequence
                                                                               0
                                                                                                   Score 18.4;
Pred. No. 1
                                                                                  Mismatches
                                                                                                   DB 1;
                                                                                                                       Length 20;
                                                                                                                                                                                                                                                                        infection by pathogenic
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                                                                               Gaps
                                                                             0;
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RESULT 751 AAF28586/c

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ARCSSILT 752
AACSSI26/C
ID AACSSI
XX AACSSI
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention relates to a composition of mammalian myeloid progenitor cells, where at least 95% of the cells are characterised as ckit (CD117 protein), IL-7Ralpha+ (interleukin-7 receptor alpha) and lin-(lineage negative). The composition is useful in transplantation, experimental evaluation, and as a source of lineage and cell specific products, including mRNA species useful in identifying genes specifically expressed in these cells, and as targets for the discovery of factors or
Crohn's disease; atopic dermatitis; autoimmune anaemia; bursitis; cholecystitis; diabetes mellitus; emphysema; atrophic gastritis; inflammatory bowel disease; multiple sclerosis; myasthenia gravis; myocardial inflammation; pericardial inflammation; osteoarthritis; osteoporosis; psoriasis; Reiter's syndrome; rheumatoid arthritis;
                                                                                                                                                     Antigen presenting cell expression protein; APEX-1; APEX-2; APEX-3; extracellular domain; immunoglobulin-like domain; Ig-like structure; N-glycosylation site; transmembrane domain; cytoplasmic domain, PCR; SH2-binding motif; asthma; arteriosclerosis; AIDS; cirrhosis; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               expressed in these cells, and as targets for the discovery of factors or molecules that can affect them. The present sequence is a PCR primer used in the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mammalian myeloid progenitor cell composition, useful for transplantation, experimental evaluation and as a source of lineage and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; myeloid progenitor cell; transplantation; CD117; IL-7Ralpha; interleukin-7 receptor alpha;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     02-APR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAC86126
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 5 A; 7 C; 3 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cell specific products.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-122949/13.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               29-JUN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200100019-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Epo-R PCR primer #1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAF28586 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                 29-AUG-2001
                                                                                                                                                                                                                                                                                                             Primer JNF14 to isolate APEX cDNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (STRD ) UNIV LELAND STANFORD JUNIOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               391
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            standard; cDNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AGTGCTGGGATTACAGGCGT 410
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Page 24; 35pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               99US-0141421P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 18.4;
Pred. No. 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PCR primer; c-kit;
Epo-R; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
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RESULT 753
AAH20704/c
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Query Match
Best Local Similarity
Matches 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                inflammation; cancer; autoimmune dis graft versus host disease; systemic polymerase chain reaction; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel Antigen presenting cell expression protein useful for treating asthma, arteriosclerosis, autoimmune diseases, AIDS, cirrhosis, Crohn's disease and atopic dermatitis.
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                                                                                                                                                                                          AAH20704 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
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                                            Antisense; phosphorothioate; human; telomeric repeat binding factor inhibitor; premature aging; hyperproliferative disorder; cancer;
                                                                                             Human telomeric repeat binding
                                                                                                                                                            AAH20704;
Homo sapiens
                              cytostatic; ss.
                                                                                                                                                                                                                                                                                                         867
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     50; Page 83; 112pp; English.
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ე
                                                                                                                                                                                                                                                                                          GGGATTACAGGCGTGAGCCA 886
                                                                                                                                                                                                                                                                          GGGATTACAGGTGTGAGCCA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                          APEX
                                                                                                                                                                                                                                                                                                                                                                                                          BP; 4 A; 8 C; 3
                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Finger J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       99US-0172025P
                                                                                                                                                                                                                                                                                                                                                                                                                                        proteins are
APEX
                                                                                                                                                                                                                                                                                                                                                       1.9%;
                                                                                                                                                                                                                                                                                                                                                                                                          G;
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                                                                                                                                                                                                                                                                                                                                         Score 18.4; D
Pred. No. 1.3e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         8
                                                                                                                                                                                                                                                                                                                                                                                                          5 T; 0 U; 0 Other;
                                                                                               factor 2 oligonucleotide 111432.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   disease; graft rejection; amplify;
mic lupus erythematosus;
                                                                                                                                                                                                                                                                                                                                                          1.3e+03
                                                                                                                                                                                                                                                                                                                                                                           DB 1;
                                                                                                                                                                                                                                                                                                                                                                         Length 20
                                                                                                                                                                                                                                                                                                                                             Indels
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RESULT 754
AAH20699/c
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                                                                                                                                                                                                                                                      Matches
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Best Local
                                                                                                                                                                                                                                                                                                                                                                      This invention describes a novel antisense compound (I) 8-30 nucleobases in length targeted to a polynucleotide encoding human telomeric repeat binding factor 2 (II) which specifically hybridizes with, and inhibits the expression of (II). (I) is useful for treating a human having a disease or condition associated with (II) such as premature aging or a hyperproliferative disorder especially cancer, by inhibiting the expression of (II) in human cells or tissues. (I) is useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Antisense compounds targeted to nucleic acid encoding telomeric repeat binding factor 2 useful for treating conditions such as premature aging
                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 5 A; 7 C; 3 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 3; Page 81; 108pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14-DEC-2000; 2000WO-US033954
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       modified_base
                                                                                                                                                                                                                                                                                                                                                  diagnostics, therapeutics, prophylaxis and as research reagents and kits. The products of the invention have cytostatic activity. This sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2001-398071/42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO200143752-A1
Homo sapiens
                       cytostatic; ss.
                                Antisense; phosphorothioate; human; telomeric inhibitor; premature aging; hyperproliferative
                                                                     Human telomeric repeat binding factor 2 oligonucleotide 111427.
                                                                                             13-AUG-2001
                                                                                                                                          AAH20699
                                                                                                                                                                                                                                                                                                                                       represents an antisense oligonucleotide used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (ISIS-) ISIS PHARM INC
                                                                                                                     AAH20699;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               diseases such
                                                                                                                                                                                                                                                                                                                              invention
                                                                                                                                                                                                                             863
                                                                                                                                                                                                                                                      al Similarity
19; Conserv
                                                                                                                                                                                                      20
                                                                                                                                                                                                                             TCCTGGGATTACAGGCGTGA 882
                                                                                                                                            standard; DNA;
                                                                                                                                                                                                        TGCTGGGATTACAGGCATGA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cowsert LM;
                                                                                                                                                                                                                                                     Conservative
                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        99US-00467642
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /*tag= c
/mod_base= OTHER
/note= "2-O-methoxyethyl"
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/mod_ba
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 as cancer.
                                                                                                                                                                                                                                                                1.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     _base= OTHER
e= "2-0-methoxyethyl"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                base=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "phosphorothioate backbone"
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                                                                                                                                                                                                                                                  0,
                                                                                                                                                                                                                                                                   Score 18.4;
Pred. No. 1
                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                 .3e+03;
                                                                                                                                                                                                                                                                               DB 1;
                                    repeat binding factor e disorder; cancer;
                                                                                                                                                                                                                                                                                                                                            illustrate
                                                                                                                                                                                                                                                                              Length 20;
                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                         0
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RESULT 755
AAS29507/c
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                                                                                                                                                                                                                                                                  Query Match
Best Local S
Matches 19
Homo sapiens
                                  Human; mdm2; hyperproliferative disorder; cancer; psoriasis; atherosclerosis; tumour; cytostatic; anti psoriatic;
                                                                                                                                                                                                                                                                                                                   Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                          in length targeted to a polynucleotide encoding human telomeric repeat binding factor 2 (II) which specifically hybridizes with, and inhibits the expression of (II). (I) is useful for treating a human having a disease or condition associated with (II) such as premature aging or a hyperproliferative disorder especially cancer, by inhibiting the expression of (II) in human cells or tissues. (I) is useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Antisense compounds targeted to nucleic acid encoding telomeric repeat binding factor 2 useful for treating conditions such as premature agin and diseases such as cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Key
modified_base
                                                                         Human mdm2 antisense oligonucleotide 31631.
                                                                                                                                                     AAS29507 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                    diagnostics, therapeutics, prophylaxis and as research reagents and kits. The products of the invention have cytostatic activity. This sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           modified_base
                                                                                                   21-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO200143752-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            modified_base
                                                                                                                                                                                                                                                                                                                                            represents an antisense oligonucleotide used to illustrate the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This invention describes a novel antisense compound (I) 8-30 nucleobases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (ISIS-) ISIS PHARM INC
                         arteriosclerotic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2001-398071/42.
                                                                                                                                                                                                                                         735
                                                                                                                                                                                                                   20
                                                                                                                                                                                                                                                                  19;
                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                         TGGGACTACAGGCGCCCACC 754
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Page 81; 108pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cowsert LM;
                                                                                                                                                                                                                                                                                                                  BP; 2 A; 7 C; 8 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                    Conservative
                                                                                                 (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    99US-00467642
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note= "
13. .20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   mod
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /*tag= a
mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        note= "phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  *tag=
                                                                                                   entry)
                                                                                                                                                                                                                                                                             1.9%;
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                       vasotropic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "2-O-methoxyethyl"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "2-O-methoxyethyl"
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                                                                                                                                                     ВÞ
                                                                                                                                                                                                                                                                  0,
                                                                                                                                                                                                                                                                              Score 18.4;
Pred. No. 1
                                                                                                                                                                                                                                                                  Mismatches
                      antisense; phosphorothioate;
                                                                                                                                                                                                                                                                                1.3e+03;
                                                                                                                                                                                                                                                                                          DB 1; Length 20;
                                                                                                                                                                                                                                                                  Indels
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                                                                                                                                                                                                                                                                  Gaps
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RESULT 756
AAS29485/c
ID AAS294
XX
AC AAS294
XX
DT 21-NOV
XX

standard;

DNA;

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21-NOV-2001 AAS29485; AAS29485

(first entry)

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                                                                                                                               cc length targeted to the 5' untranslated region, translation termination cc codon region, 3' untranslated region, coding region or translation start csite of a nucleic acid encoding human mdm2, where the antisense compound cc modulates the expression of human mdm2. The antisense oligonucleotides of the invention are useful for encoding human mdm2 and for inhibiting the expression of human mdm2. They may be used for treating an animal having cd disease or condition associated with amplification of mdm2 gene or overexpression of mdm2 e.g. a hyperproliferative disorder such as cancer (blood, brain, breast, lung, or a soft tissue cancer) and psoriasis, cf ibrosis, atherosclerosis or restenosis, tumours, colorectal carcinoma cd and chronic myelogenous leukemia. The antisense compound may be cardisense compound reduces hyperproliferation of human cells. The method, which involves the use of the antisense compound, is also useful cor detecting the role of mdm2 expression in various cell functions and chysiological processes and useful in both clinical research and conjugant tools. AAS29242-AAS29507 represent the human mdm2 antisense
                                                                               Query Match
Best Local (
                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Key
modified_base
                                                                                                                             Sequence 20 BP; 3 A; 6 C; 5 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to antisense compounds, 8-30 nucleobases length targeted to the 5' untranslated region, translation termination
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  of a nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                An antisense compound, useful for treating e.g. cancer, comprises nucleobases targeted a region (e.g. translation termination codon region)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Miraglia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 9; Page 18; 81pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2001-535565/59.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (NERO/)
(GRAH/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               26-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                02-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US2001016575-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  26-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (MIRA/) MIRAGLIA L J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MONI/
                                868 GGATTACAGGCGTGAGCCAC 887
20
                                                               l Similarity
19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MONIA B P.
COWSERT L M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NERO P.
GRAHAM M J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ę
                                                                 Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nero P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                98US-00048810.
99US-00280805.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /mod_base= OTHER
/not== "OTHER= All phosphorothioate linkages,
/note= "OTHER= All phosphorothioate linkages,
additionally bases 1-6 and bases 15-20 are 2'-O-
methoxyethyl bases, and bases 7-14 are deoxynucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /*tag= a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                encoding
                                                                             1.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Graham MJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  human
                                                               <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               for mdm2.
                                                                               Score 18.4; DB 1;
Pred. No. 1.3e+03;
                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cowsert LM;
                                                                                              Length 20;
                                                               0,
                                                               Gaps
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Query Match
Best Local
                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to antisense compounds, 8-30 nucleobases in length targeted to the 5' untranslated region, translation codon region, 3' untranslated region, coding region or translation start site of a nucleic acid encoding human mdm2, where the antisense compound modulates the expression of human mdm2. The antisense oligonuclectides of the invention are useful for encoding human mdm2 and for inhibiting the expression of human mdm2. They may be used for treating an animal having a disease or condition associated with amplification of mdm2 gene or overexpression of mdm2 e.g. a hyperproliferative disorder such as cancer (blood, brain, breast, lung, or a soft tissue cancer) and psoriasis, fibrosis, atheroselerosis or restenosis, tunours, colorectal carcinoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    An antisense compound, useful for treating e.g. nucleobases targeted a region (e.g. translation of a nucleic acid encoding human mdm2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human
                                                                                                                                                                                                                                                   and chronic myelogenous leukemia. The antisense compound may be administered with a chemotherapeutic agent to overcome drug resistance. The antisense compound reduces hyperproliferation of human cells. The method, which involves the use of the antisense compound, is also useful for detecting the role of mdm2 expression in various cell functions and physiological processes and useful in both clinical research and diagnostic tools. AAS29242-AAS29507 represent the human mdm2 antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26-MAR-1998;
26-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           atherosclerosi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 4; Page 18; 81pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-535565/59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Miraglia LJ, Nero P,
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(GRAH/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   02-JAN-2001; 2001US-00752983
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                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                  oligonucleotides of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (/INOM)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (MIRA/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cosclerosis; tumour; arteriosclerotic; va
                                            899
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        mdm2; hyperproliferative disorder; cancer; psoriasis; sclerosis; tumour; cytostatic; anti psoriatic;
20
                                                                                        l Similarity
19; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NERO
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COWSERT L M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GRAHAM M J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MIRAGLIA L
                           TCTTGGCTCACTGCAACCTC 687
                                                                                                                                                                                   20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  antisense oligonucleotide
                                                                                                                                                                                   BP;
                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         יסי.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              98US-00048810.
99US-00280805.
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additionally bases 1-6 and bases 15-20 are 2'-0-
methoxyethyl bases, and bases 7-14 are deoxynucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  mod_base= OTHER
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                                                                                                                                                                                   A
                                                                                                                                                                                                                             the present
                                                                                                              1.9%;
                                                                                                                                                                                   4 C; 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       vasotropic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Graham MJ,
                                                                                        0,
                                                                                                                                                                                   G; 3 T; 0 U; 0 Other;
                                                                                        Score 18.4; D
Pred. No. 1.3e
0; Mismatches
                                                                                                                                                                                                                                  invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       antisense;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    31468
                                                                                                                 .3e+03;
                                                                                                                                            DB
                                                                                                                                         1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cowsert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       phosphorothicate; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cancer, comprises
termination codon region)
                                                                                                                                       Length
                                                                                             Indels
                                                                                                                                         20;
                                                                                                                                                                                                                                                                                                                                is also useful
                                                                                             0
                                                                                           Gaps
                                                                                             0;
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RESULT 757

밁 8

Matches Query Match Best Local

Similarity

95.0%; 1.9%;

Conservative

0; Mismatches Score 18.4; DB 1 Pred. No. 1.3e+03

DB 1;

Length Indels

20,

٥,

Gaps

0

385 19;

TCCCAAAGTGCTGGGATTAC 404

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The present invention relates to a polynucleotide encoding polymorphic CC Kinase anchor protein (AKAP), with isoleucine residue at position 646 CC replaced with valine, leucine or phenylalanine. AKAP is useful for CC detecting an allelic variant of a human AKAP10 gene which is indicative of an alteration in signal transduction, where the alteration is related CC neurological, neurodegenerative disorders, obesity, diabetes and CC peripheral retinopathies, especially the disorders including Alaheimer's CC disorder and retinitis pigmentosa. The method of the invention is useful for indicating susceptibility to morbidity and/or increased or early CC comprises a nucleotide other than A at position T corresponding to 2073 of AKAP, or a polymorphic region of AKAP10 CC comprises a nucleotide other than A at position T corresponding to position 2073 of AKAP, or a polymorphic region of AKAP10 CC comprises and single-stranded conformation polymorphism analysis, or the detecting step is performed by allele specific hybridisation, primer cc specific extension, oligonucleotide ligation assay, restriction enzyme site analysis and single-stranded conformation polymorphism analysis, or the detection is by detecting a signal group from radioisotopes, enzymes, antipodes, spectrophotometric reagents, chemiluminescent corresponts, fluorescent reagents and other light producing reagents. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New polynucleotide encoding polymorphic A-Kinase anchor proteins for detecting an allelic variant of the human gene which is indicative of alteration in signal transduction, and is related to a disorder e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     bipolar; cardiovascular; cardiac; proliferative; neurodegenerative; cardiomyopathy; peripheral retinopathy; obesity; signal transduction left ventricular function; Alzheimer's disease; retinitis pigmentos
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; polymorphic A-Kinase anchor protein; AKAP; disorder; neurological;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Braun
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10-JUL-2000; 2000US-0217251P.
13-OCT-2000; 2000US-0240335P.
12-APR-2001; 2001US-00834700.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Target specific PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAD28754;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAD28754 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-154919/20.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO200204489-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
 Sequence
                                    present sequence is a target specific PCR primer used AKAP10-1 target sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 3; Page 90; 290pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (SEQU-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENOM INC
   20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PCR
   BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         primer; ss.
   6 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   primer #2 to amplify human AKAP10-1 target sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         entry)
   5 C; 4 G; 5 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20
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     0 U;
       0 Other
                                                             ö
                                                           amplify human
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RESULT 758
RAL40357
ID AAL403
XX AAL403
XX AAL403
XX Human
XX Human
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KW Antill
KW AAL401
KW AAL403
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ABK68939
      BXXXB
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Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to an antisense oligonucleotide compound of 8 to 50 nucleotides in length that is targeted to a nucleic acid molecule encoding caspase 6, where the oligonucleotide specifically hybridises with and inhibits the expression of caspase 6. The oligonucleotide of the invention specifically hybridises to and inhibits expression of caspase 6 in cells or tissues. The oligonucleotides can be administered caspase 6 condition associated with caspase 6, such as Rieger's syndrome or ataxia telangiectasia, hyperproliferative disorder, a haematopoietic disorder, a bone metabolism or cholesterol disorder, a haematopoietic cancer, neurological conditions such as Rieger's disease and other regulated apoptotic pathological conditions. This polynucleotide sequence represents a human caspase 6 oligonucleotide relating to the invention.

NOTE: This phosphorothioate oligonucleotide sequence has 2'-MOE wings and
   02-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Muscular; cytostatic; nootropic; neuroprotective; ophthalmological; antilipaemic; osteopathic; caspase 6; Rieger's syndrome; bone metabolism; ataxia telangiectasia; hyperproliferative disorder; cholesterol disorder; haematopoietic disorder; cancer; neurological; Alzheimer's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human caspase
                                                               ABK68939
                                                                                                                            ABK68939 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 3; Page 89; 141pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                An antisense oligonucleotide of 8 to 50 nucleotides in length that inhibits caspase 6, is useful for treating Rieger's syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Brown-Driver VL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               04-OCT-2000; 2000US-00679299
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        apoptotic; human; ds.
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
                                                                                                                                                                                                                                                                                                                                           545
                                                                                                                                                                                                                                                                                  سر
                                                                                                                                                                                                                                                                                                               AGCCTCCCAAGTAGCTGGGA 564
                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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(first entry).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Zhang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; 5 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                           1.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ħ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20
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                                                                                                                                                                                                                                                                                                                                                                                                     <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Watt AT;
                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 18.4;
Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                     1.3e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 1; Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                     0
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                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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RESULT 760 AAL38181/c

AAL38181 standard; DNA; 20 BP

29-AUG-2003

(revised)

AAL38181;

밁 Ş

0

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modulating the expression of human phosphorylase kinase beta. The CC antisense compounds, particularly antisense oligonucleotides, target and cinhibit the expression of human phosphorylase kinase beta. The antisense compounds are useful for inhibiting the expression of human phosphorylase kinase beta in human cells or tissues and for treating an animal, CC particularly a human suspected of having or being prone to a disease or condition associated with expression of phosphorylase kinase beta such as a metabolic disorder e.g. diabetes. The compounds are useful for CC diagnostics, therapeutics and as research reagent, e.g. prophylactically to prevent or delay infection, inflammation or tumour formation. The CC antisense compounds are useful in the preparation of a pharmaceutical CC formulation. They are highly specific, have an enhanced affinity for the nucleic acid target, and are safely and effectively administered to humans. ABK68888-ABK68965 represent human phosphorylase kinase beta
                                                                                                                            Query Match
Best Local S:
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel antisense oligonucleotide which inhibits expression of phosphorylase kinase beta, useful for treating metabolic disorder e.g. diabetes, prevent or delay infection, inflammation or tumor formation.
                                                                                                                                                                                                                                                               Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           antiinflammatory; cytostatic; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human phosphorylase kinase beta antisense oligonucleotide #52.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 3; Page 83; 132pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         12-SEP-2001; 2001WO-US028586
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200222637-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 nfection;
                                                                                                                        Local Similarity hes 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    present invention relates to antisense compounds and
                                                              636 TCTGTCACCCAGGCTGGAGT 655
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          phosphorylase kinase beta; metabolic disorder; diabetes;
ion; inflammation; tumour formation; antidiabetic;
L
                                                                                                                                   Conservative
                                                                                                                                                                                                                                                               BP; 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2000US-00662250.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /mod_base= OTHER
/note= "OTHER= Phosphorothioate internucleotide linkages,
optionally bases 1-5 and 16-20 are 2'-methoxyethoxy (2'-
MOE) bases, where the 2'-MOE cytidines are also
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                               A; 6 C; 6 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    methylcytidines"
                                                                                                                                                              1.9%;
95.0%;
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                                                                                                                                                           Score 18.4;
Pred. No. 1.
20
                                                                                                                                   Mismatches
                                                                                                                                                           1.3e+03
                                                                                                                                                                                                 DB 1;
                                                                                                                                                                                          Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       methods for
                                                                                                                                   0
                                                                                                                                   Gaps
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15-AUG-2002

(first entry)

Hepatotrophic;

immunomodulatory; cytostatic; antiinflammatory; hepatitis;

domain death mRNA agonist inhibitor SEQ

ij

24

BH3

interacting

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RESULT 761
AAL38190
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                                                                                                                                                                                                                                                                                                                                     The invention relates to a compound 8 to 50 nucleotides in length CC targeted to a nucleic acid molecule encoding a BH3 interacting domain CC death agonist, where the compound specifically hybridises with and CC inhibits the expression of the BH3 interacting domain death agonist. The CC compound of the invention is useful for inhibiting the expression of the CC BH3 interacting domain death agonist in cells or tissues. The compound is CC also useful for treating an animal having a disease or condition CC associated with the BH3 interacting domain death agonist, e.g. CC disorder, immunological disorder, or a disease or condition of the liver CC e.g., hepatitis, or a condition associated with apoptosis. The compound CC is useful for diagnostice, therapeutics, prophylaxis and as research CC reagents and kits. This polynucleotide sequence represents an antisense CC oligonucleotide inhibitor of the DNA from human BH3 interacting domain CC death agonist RNA of the invention. NOTE: This sequence is a chimeric CC oligonucleotide inhibitor of the DNA from human BH3 interacting domain CC oligonucleotide inhibitor of the DNA from human BH3 interacting domain CC oligonucleotide inhibitor. The wings are composed of 2-methoxyethyl (2' hyber nucleotides. The internucleoside (backbone) linkages are pauges are composed of 2-methoxyethyl (2' phosphorothicate (P=S) throughout the oligonucleotide. (Updated on 29-AUG phosphorothicate (P=S) throughout the oligonucleotide.
                                                                                                                                                                                                                                  Query Match
Best Local
                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel antisense compound targeted to nucleic acid molecule encoding the BH3 interacting domain death agonist, useful for treating animals with diseases associated with BH3 interacting domain death agonist, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            haemostatic; BH3 interacting domain death agonist; liver disease; haematopoietic disorder; developmental disorder; immunological disorder; hyperproliferative disorder; apoptosis; human; chimeric; 2'-methoxyethyl; 2'-MOE; phosphorothioate backbone; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-SEP-2000; 2000US-00657346.
07-MAR-2001; 2001US-00800631.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 3; Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                31-AUG-2001; 2001WO-US027316.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200220547-A1.
               AAL38190
                                                     AAL38190 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2002-393838/42
                                                                                                                                                                             538
                                                                                                                                                                                                                                                                                                                            ç
                                                                                                                                            20
                                                                                                                                                                                                                 l Similarity
19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SISI
                                                                                                                                                                           CTGCCTCAGCCTCCCAAGTA 557
                                                                                                                                                                                                                                                                                           20
                                                                                                                                                                                                                                                                                                                            standardise OS field)
                                                                                                                                            CTGCCTCAGCCTCCCGAGTA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Wyatt JR;
                                                                                                                                                                                                                                                                                       BP; 4 A; 4 C; 9 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PHARM INC.
                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               86; 171pp; English
                                                                                                                                                                                                                                1.9%;
                                                                                                                                                                                                                 0
                                                                                                                                                                                                                                  Score 18.4;
Pred. No. 1.
                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                  .3e+03
                                                                                                                                                                                                                                                     DB 1;
                                                                                                                                                                                                                                                   Length
                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                       20;
                                                                                                                                                                                                                   <u>.</u>.
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RESULT 762
ABQ74794/c
ID ABQ747
XX
AC ABQ747

ABQ74794 standard; DNA;

20

ВP

ABQ74794;

В δ

H

TCTCGGCTCACTACAACCTC TCTCGGCTCACTGCAACCTC 987

20

0

896

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The invention relates to a compound 8 to 50 nucleotides in length CC targeted to a nucleic acid molecule encoding a BH3 interacting domain CC death agonist, where the compound specifically hybridises with and CC inhibits the expression of the BH3 interacting domain death agonist. The CC compound of the invention is useful for inhibiting the expression of the BH3 interacting domain death agonist. The CC also useful for treating an animal having a disease or condition CC associated with the BH3 interacting domain death agonist, e.g. CC disorder, immunological disorder, or a disease or condition CC is useful for diagnostics, therapeutics, prophylaxis and as research CC e.g., hepastitis, or a condition associated with apoptosis. The compound CC is useful for diagnostics, therapeutics, prophylaxis and as research CC eath agonist RNA of the invention. NOTE: This sequence is a chimeric CC diagnocleotide inhibitor of the DNA from human BH3 interacting domain CC death agonist RNA of the invention. NOTE: This sequence is a chimeric CC by five-nucleotides. The intertucleoside is a chimeric CC by five-nucleotides. The intertucleoside is a chimeric CC by five-nucleotides. The intertucleoside (backbone) linkages are CC phosphorothioate (P=S) throughout the oligonucleotide. (Updated on 29-AUG cc. 2003 to standardise OS field)
Matches
                                Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       haemostatīc; BH3 interacting domain death agonist; liver disease; haematopoietic disorder; developmental disorder; immunological disorder; hyperproliferative disorder; apoptosis; human; chimeric; 2'-methoxyethyl; 2'-MOS; phosphorothicare harrhome. As
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel antisense compound targeted to nucleic acid molecule encoding the BH3 interacting domain death agonist, useful for treating animals with diseases associated with BH3 interacting domain death agonist, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    07-SEP-2000; 2000US-00657346.
07-MAR-2001; 2001US-00800631.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  29-AUG-2003
15-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  31-AUG-2001; 2001WO-US027316
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chimeric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 3; Page 86; 171pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-393838/42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Zhang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     14-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200220547-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens.
                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BH3 interacting domain death mRNA agonist inhibitor SEQ
   19;
                                Similarity
                                                                                                                          20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Wyatt
   Conservative
                                                                                                                          BP; 4 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (revised)
                                                                                                                          9 C;
                                1.9%;
                                                                                                                          2 G; 5 T;
   0
                                Score 18.4;
Pred. No. 1.
      Mismatches
                                                                                                                          0 U; 0 Other
                                .3e+03;
                                                                DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 antiinflammatory;
                                                             Length
         Indels
         <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  IJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 hepatitis;
         Gaps
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RESULT 763
ACC55324/c
ID ACC553
XX
AC ACC553
XX
DT 27-JUN
XX
DE Human
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                                                                                                                              Query Match
Best Local S
Matches 19
                                                                                                                                                                             The present invention describes compounds of 8-30 nucleobases antisense to a nucleic acid encoding human or mouse tumour necrosis factor receptor 2 (TNRR2). Also described is a method for inhibiting expression of human or mouse TNFR2 comprising contacting cells or tissues in vitro with one of the claimed compounds. The antisense compounds are used to treat a disease or condition associated with expression of TNFR2. The present sequence represents a human TNFR2 antisense chimeric phosphorothicate oligonucleotide, which is given in the present invention
Human ADAMTS13 STS marker GL4-1 5'
                                                                                                                                                                                                                                                                                    New compounds antisense to nucleic acid encoding human necrosis factor receptor 2 are useful to treat disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Tumour necrosis factor receptor 2; TNFR2; antisense oligonucleotide; phosphorothioate; 2'-O-methoxyethyl; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   24-OCT-2002
                                   ACC55324;
                                                   ACC55324 standard;
                                                                                                                                                                 Sequence 20
                                                                                                                                                                                                                                                           Example 15; Col 47; 69pp; English.
                                                                                                                                                                                                                                                                             mouse tumor necrosis factor receptor 2 expression.
                                                                                                                                                                                                                                                                                                                               Bennett CF,
                                                                                                                                                                                                                                                                                                                                                                27-APR-2001; 2001US-00844634.
                                                                                                                                                                                                                                                                                                                                                                                 27-APR-2001; 2001US-00844634.
                                                                                                                                                                                                                                                                                                                                                                                                   25-JUN-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                    US6410324-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                           modified_base
                  27-JUN-2003
                                                                                                                                                                                                                                                                                                                                                (ISIS-)
                                                                                                                                                                                                                                                                                                               2002-606814/65.
                                                                                                              546
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TNFR2
                                                                                             20
                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                ISIS PHARM INC.
                                                                                                              GCCTCCCAAGTAGCTGGGAC 565
                                                                                             GCCTCCCAAGTAGATGGGAC 1
                                                                                                                                                                BP; 3 A; 6 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                               Watt AT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note= "
16. .20
                                                                                                                                                                                                                                                                                                                                                                                                                                                             /mod_base= OTHEK
/note= "2'-O-methoxyethyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               mod
d
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                    note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                   *tag=
                                                    DNA;
                                                                                                                                       1.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               _base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                            base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   oligonucleotide SEQ ID NO:44
                                                                                                                                                                                                                                                                                                                                                                                                                                    "2'-O-methoxyethyl nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "phosphorothioate linkages"
                                                    20
                                                    ВP
                                                                                                                               0
                                                                                                                                       Score 18.4; DB 1;
Pred. No. 1.3e+03;
                                                                                                                               Mismatches
 PCR primer.
                                                                                                                                                DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     nucleotides"
                                                                                                                                              Length 20;
                                                                                                                               Indels
                                                                                                                                                                                                                                                                                     or mouse tumor associated with
                                                                                                                               0
                                                                                                                               Gaps
                                                                                                                               0
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RESULT 764
ABZ71056/c
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to a novel method for identifying subjects at risk of developing thrombotic thrombotytopenic purpura (TTP) disease, comprising providing nucleic acid having a disintegrin and metalloproteinase containing thrombospondin 1-like domains 13 (ADAMTS13) gene from a subject, and detecting the presence or absence of one or more variations in the ADAMTS13 gene. The method of the invention has thrombolytic and haemostatic activity. The methods and compositions of the present invention are useful for the diagnosis and treatment of, and/or analysing risks for thrombotic thrombocytopenic purpura. The present sequence is used in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; thrombotic thrombocytopenic purpura; TTP; disintegrin; metalloproteinase; thrombospondin 1-like domains 13; ADAMTS13; thrombolytic; haemostatic; PCR; primer; RT-PCR; 5' RACE; 3' RACE;
                                                                                                                                                                                                                             Human; HKR1; cytostatic; HKR1 inhibitor; hyperproliferative disorder; cancer; antisense oligonucleotide; 2'-O-methoxyethyl; 2'-MOE; control; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Identifying risk of developing thrombotic thrombocytopenic disease, using a novel disintegrin and metalloproteinase conthrombospondin 1-like domains genes and proteases.
                   modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABZ71056 standard; DNA;
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16-AUG-2002;
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modified_base
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                                                                                                                                                                                 Homo sapiens
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Pred. No. 1.3e+03;
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RESULT 765
ADA20921/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New compounds, particularly antisense oligonucleotides targeted to a nucleic acid encoding HKR1, useful for treating a disease/condition associated with HKR1, such as hyperproliferative disorder, e.g. lung,
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                                     Human BAX chimeric phosphorothioate oligonucleotide
                                                                                                                                                                  ADA20921 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
BCL2-associated X;
                                                                                  20-NOV-2003
                                                                                                                            ADA20921;
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19; Conserv
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/mod_base= OTHER
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BAX; nootropic;
                                                                                                                                                                                                                                                                                                                                                                                  1.9%;
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Pred. No. 1.3e+03;
                                                                                                                                                                                                                                                                                                                                                                Mismatches
  neuroprotective;
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antiparkinsonian;
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The present invention describes a compound (I) 8-50 nucleobases in length carageted to a nucleic acid molecule encoding BCL2-associated X (BAX) CC protein, where the compound specifically hybridises with the nucleic acid compound specifically hybridises with the nucleic acid compound specifically hybridises with at least 8-nucleobase portion cof an active site on a nucleic acid molecule encoding BAX protein. Also compound specifically hybridises with at least 8-nucleobase portion cof an active site on a nucleic acid molecule encoding BAX protein. Also composition comprising (I) and a pharmaceutical carrier crisuses comprising contacting the expression of BAX protein in cells or tissues comprising contacting the cells or tissues with (I); and (3) cc treating an animal having a disease or condition associated with BAX protein is inhibited. (I) has nootropic, neuroprotective, contribution associated with BAX protein, and can be used in antisense therapy, and as a BAX crisusesion of BAX protein, and for treating a disease or condition cassociated with BAX protein, and for treating a disease, Hodgkin's disease, activation and some compounds (I) are useful for modulating the content of BAX protein, and for treating a disease, Hodgkin's disease, activation is disease, activation and some compounds (I) and sease, activation and sease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Zhang
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present sequence represents a human B
oligonucleotide, which is used in an
                                                                                                                               scrapie infection, or a condition that arises from aberrant apoptosis. The compounds are useful as research reagents and in diagnostics. The
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                                                                      sequence represents a human BAX chimeric phosphorothioate
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/note= "2'-O-methoxyethyls"
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/note= "2'-O-methoxyethyls"
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Query Match

Sequence 20

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ACF39682
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Matches 19; Conserv
The present invention describes a compound (I) that is 8-50 nucleobases in length: (a) targets a nucleic acid molecule encoding major histocompartibility complex (MHC) class II transactivator, and specifically hybridises with the nucleic acid encoding the MHC class II transactivator, and inhibits the expression of MHC class II transactivator, or (b) specifically hybridises with at least an 8-nucleobase portion of an active site on a nucleic acid molecule encoding
                                                                                                                                      New antisense oligonucleotides for modulating MHC class II transactivator gene expression, particularly useful for treating autoimmune disorders such as transplant rejection, Alzheimer's disease, or multiple sclerosis,
                                                                                                                                                                                                                                                                                                  04-DEC-2002; 2002WO-US038616
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            nootropic; neuroprotective; immunostimulant; autoimmune disort MHC Class II transactivator inhibitor; infection; transplant; diabetes; rheumatoid arthritis; cancer; Alzhelmer's disease; multiple sclerosis; severe combined immunodeficiency disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; major histocompatibility complex class II transactivator; MHC class II transactivator; antisense modulation; immunosuppres antinicrobial; antidiabetic; antirheumatic; antiarthritic; cytos
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MHC class II transactivator antisense oligonucleotide SEQ ID NO:85
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are 5-methylcytidines"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    transplant rejection;
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RESULT 767
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cc antidiabetic, antirheumatic, antiarthritic, cytostatic, nootropic, concurportective and immunostimulant activities, and can be used as an MHC class II transactivator inhibitor. The MHC class II transactivator condition associated with MHC class II transactivator condition associated with MHC class II transactivator, e.g. condition associated with MHC class II transactivator in condition associated with MHC class II transactivator in condition condition associated with MHC class II transactivator in condition condition associated with MHC class II transactivator in collision of the expression of MHC class II transactivator in collision conditions. In particular, these diseases include transplant conjugate in the condition of the expression of of 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local
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                          This invention relates to a novel antisense compounds that modulate expression of oestrogen receptor beta (ERSB). Oestrogen is a steroid hormone that exerts a wide range of effects throughout the human bod being primarily involved in female sexual maturation. Additionally, however, oestrogen targets male reproductive tissues, is known to be important in bone maintenance and plays a protective role in the cardiovascular system. This hormone receptor, ERSB (also known as ER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
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                                                                                                                                                                                                                                                                                                                                                                                             or lymphoma.
                                                                                                                                                                                                                                                                                                                                                                                                                       New antisense oligonucleotides for modulating estrogen receptor beta gene expression, particularly useful for treating cancers, specifically leiomyoma, pancreatic cancer, prostate cancer, breast cancer, bone cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Dobie KW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  07-DEC-2001; 2001US-00005058
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   bone maintenance; cardiovascular system; ER beta; o
ERS2; Alzheimer's; uterine leiomyomata; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Oestrogen receptor beta; ERSB; steroid hormone;
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                                                                                                                                                                                                                                                                                                                         Claim 3;
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19; Conserv
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Pred. No. 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              female sexual
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length
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c; kidney neoplasm; ss;
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RESULT 768
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
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Matches 19
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                                                     The invention comprises antisense oligonucleotides which are targeted to the human mdm2 gene. The antisense oligonucleotides of the invention are useful for reducing hyperproliferation of human cells. The antisense oligonucleotides are also useful for treating: hyperproliferative disorders (e.g. cancer), psoriasis, fibrosis, atherosclerosis, or restenosis. The antisense oligonucleotides are also useful for modulating apoptosis, and for increasing expression of p21. The present DNA sequence represents a human mdm2 gene antisense oligonucleotide of the invention. The present sequence contains 2'-methoxverthoxv-residence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          antisense oligonucleotide; human; mdm2; hyperproliferation; hyperproliferative disorder; cancer; psoriasis; fibrosis; atherosclerosis; restenosis; apoptosis modulation; p21; ss; 2'-methoxy-residue; phosphorothioate backbone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADD21681 standard; DNA; 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel antisense compound targeted to 5' untranslated region, region, or intron:exon junction of nucleic acid molecule encouseful for treating e.g. cancer, psoriasis or restenosis by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human mdm2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Miraglia LJ,
Manoharan M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         04-DEC-2001; 2001US-00005344
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                Claim 4;
                                                                                                                                                                                                                                                                                                                                                                                                                        mdm2 expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2003-577263/54.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        02-DEC-2002; 2002WO-US038281
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                                                                                                                                                                                                                                                                                                                                                             SEQ ID NO 246; 289pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 antisense oligonucleotide #244.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Graham MJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ВÞ
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Pred. No. 1.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Koller
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by inhibiting
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    coding
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Matches 19
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Best Local Similarity
Matches 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 6 A; 4 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                       the human mdm2 gene. The antisense oligonucleotides of the invention are useful for reducing hyperproliferation of human cells. The antisense oligonucleotides are also useful for treating: hyperproliferative disorders (e.g. cancer), psoriasis, fibrosis, atherosclerosis, or restenosis. The antisense oligonucleotides are also useful for modulating apoptosis, and for increasing expression of p21. The present DNA sequence represents a human mdm2 gene antisense oligonucleotide of the invention. The present sequence contains 2'-methoxyethoxy-residues and has a
                                                                                                                                                                                                                                                                                 Novel antisense compound targeted to 5' untranslated region, region, or intron:exon junction of nucleic acid molecule encuseful for treating e.g. cancer, psoriasis or restenosis by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       antisense oligonucleotide; human; mdm2; hyperproliferation; hyperproliferative disorder; cancer; psoriasis; fibrosis; atherosclerosis; restenosis; apoptosis modulation; p21; ss; 2'-methoxyethoxy-residue; phosphorothioate backbone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human mdm2 antisense oligonucleotide #266.
                                                                                                                                                                                                                                                                                                                                                            Miraglia LJ,
Manoharan M;
                                                                                                                                                                                                                                                                                                                                                                                                                     04-DEC-2001;
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                                                                                         Sequence
                                                                                                                                                                                                                       The invention comprises antisense oligonucleotides which are targeted
                                                                                                                                                                                                                                                  Claim 4; SEQ ID NO 268; 289pp;
                                                                                                                                                                                                                                                                                                                                   WPI; 2003-577263/54.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12-JUN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO2003048315-A2
                                                                                                                phosphorothioate backbone.
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                      GGATTACAGGCGTGAGCCAC 887
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                                               Conservative
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                                                                                          6 C; 5 G;
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Pred. No. 1.
                                                          Score 18.4;
Pred. No. 1
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                                                                                                                                                                                                                                                 English.
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                                               Mismatches
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by inhibiting
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PF 25-OCT

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PR 25-OCT

RR 08-NOV

PR 09-NOV

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                                                                                                                                                                                                                                                                                  CC predisposition for or the occurrence of neurodegenerative disease in a cc subject. The method comprises detecting in a target nucleic acid obtained from the subject the presence or absence of an allelic variant of one or more polymorphic regions of one or more genes selected from uPA (Urokinase plasminogen activator), SNCG (gamma-synuclein), IDE (insulince degrading enzyme), KNSL1 (Kinesin-like protein 1), LIPA (lysosomal acid lypase), and TNFRSF6 (Tumour Necrosis Factor Receptor-SF6), where the presence of at least one of the allelic variant of one or more cocurrence of neurodegenerative disease. The genes are all located on chromosome 10. M1 is useful for determining a predisposition for or the occurrence of, and for treating neurodegenerative disease, particularly clinthemetrs disease. The present sequence is a PCR primer, which was used in the method of the invention
                                                                                                      Query Match
Best Local S
Matches 19
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08-NOV-2001; 2001US-0336929P.
08-NOV-2001; 2001US-0338010P.
09-NOV-2001; 2001US-0338363P.
04-DEC-2001; 2001US-0337052P.
28-MAR-2002; 2002US-0368919P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Determining a predisposition for or the occurrence of neurodegenerative disease, e.g. Alzheimer's disease by detecting in a target nucleic acid the presence or absence of an allelic variant of one or more polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           03-JUL-2003.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Neurodegenerative disease; uPA; SNCG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human KNSL1 sequencing primer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           29-JAN-2004
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                                                                                                                                                                                                              Sequence 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Saunders
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                                                                                                                                                                                                                                                                          method of the invention.
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                                                                                                        Similarity 95.0
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                                             CGCTAATTTTTGTATTTTCA 1083
CGCTAATTTTTGTATTTTTA 1
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Mullin KM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          disease; neuroprotective;
l0; PCR; primer; ss.
                                                                                                                                                                                                                 BP; 12 A; 2 C;
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Sampson AJ,
                                                                                                                                                                                                                    2 G; 4 T;
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Pred.
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                                                                                                           Mismatches
                                                                                                                                     18.4; DB 1
No. 1.3e+03
                                                                                                                                                                                                                 0 U; 0 Other;
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Blacker DL;
                                                                                                                                                           DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Tanzi RE,
                                                                                                                                                         Length 20;
                                                                                                           Indels
                                                                                                        0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bertram
                                                                                                        Gaps
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                                                                                                        0
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RESULT 771 ADE86781/c ID ADE867

ADE86781 standard; DNA; 20

ВP

Location/Qualifiers

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RESULT 772
AD161628
ID AD1616
XX AD1616
XX AD1616
XX AD1616
XX AD1616
XX C2-APR
XX AD1616
XX CYCOST
XX CYCOST
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XX J
OS HOMO 8
SX Synthe
XX Synthe
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 XAXFE X BXX BXX BXX BX F X F X F X F X F X B X S X S S S S S S S X S X S S X S S X S S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S X S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This sequence represents a primer which was used in the molecular phenotyping of brain and lung derived CD31+ cells. This primer was used to isolate cells which are CD31+ CD34+ CD45- CD105+ c-kit-lin-. The isolated cells were used in the composition of the invention which comprises fewer than 20% of lineage committed cells. The composition of the invention is useful for reconstituting haematopoiesis, and therefore in promoting survival, proliferation and/or differentiation of haematopoietic stem cells which may be used in treating cancers (e.g. leukaemia) or in bone marrow transplantation as well as transplantation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A composition for promoting survival, proliferation and/or differentiation of hematopoietic stem cells useful in e.g. bone marrow transplantation, comprises lin cells that express CD31, CD34 and CD105.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ss; primer; molecular phenotyping; brain; lung; CD31+ cell; lineage committed cell; survival; proliferation; differentiation; haematopoletic stem cell; cancer; leukaemia; bone marrow;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         of other organs in association with the transplantation of bone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GATA primer
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                                                          Synthetic
                                                                                   Homo sapiens
                                                                                                                                                                                cytostatic;
                                                                                                                                                                                                                                        Human SAP-1 gene targeted oligonucleotide ISIS 176450.
                                                                                                                                                                                                                                                                                                     22-APR-2004
                                                                                                                                                                                                                                                                                                                                                                ADI61628;
                                                                                                                                                                                                                                                                                                                                                                                                                        ADI61628
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 1; Page 27; 54pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-854104/79.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Fleming
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16-APR-2002; 2002US-0373127P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15-APR-2003; 2003WO-US011649
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     transplantation.
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                                                                                                                                                 ŞAP-1; splice
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (UYOR-) UNIV OREGON HEALTH SCI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WH,
                                                                                                                                                                                                                                                                                                                                                                                                                        standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AGTGCTGGGATTACAGGCAT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AGTGCTGGGATTACAGGCGT 410
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ľ
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                                                                                                                                                    antisense therapy; Serum response factor Accessory Protein;
se variant; hyperproliferative disorder; cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                     (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                           DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1.9%;
95.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 18.4;
Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
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RESULT 773
ABZ97965
ID ABZ979
XX
AC ABZ979
XX
DT 17-OCT
XX
DE Human
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                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to a compound 8-50 nucleobases in length targeted to a nucleic acid encoding Serum response factor Accessory Protein (SAP)-1 or a splice variant of SAP-1, where the compound specifically hybridizes with and inhibits the expression of SAP-1, a splice variant of SAP-1, or a truncated form of SAP-1, or hybridizes with an 8-nucleobase portion of an active site on a nucleic acid encoding SAP-1. The compound is used for inhibiting the expression of SAP-1 in cells or tissues, and for treating an animal having a disease or condition associated with SAP-1. The disease or disorder is a hyperproliferative disorder, such as breast, prostate or hematopoietic cancer. The antissues compounds may be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             modified_base
                                                                                                                                                                                                                                                                                                                      used as research reagents and diagnostics, as therapeutics, prophylaxis, to elucidate the function of particular genes, to distinguish between functions of various members of biological pathway, and in treatment regimes of cells, tissues, and animals. This sequence corresponds to an oligonucleotide of the invention which is targeted to the human SAP-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       response factor Accessory Protein (SAP)-1, for inhibiting the express of SAP-1 and for treating a disease or condition associated with the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New antisense oligonucleotides targeted to nucleic acid encoding Serum response factor Accessory Protein (SAP)-1, for inhibiting the expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 15; SEQ ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          expression of SAP-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Baker BF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-AUG-2001; 2001US-00920759.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    31-JUL-2002; 2002WO-US024369
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO2003011888-A1
                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13-FEB-2003
                                                           ABZ97965
                              17-OCT-2003
                                                                                        ABZ97965 standard;
                                                                                                                                                                                                                                     Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2003-248144/24.
                                                                                                                                                                                          665
                                                                                                                                                                                                                      19;
                                                                                                                                                             _
                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                              20
                                                                                                                                                                                          CAATCTTGGCTCACTGCAAC 684
                                                                                                                                                             CAATCTTGGCTCACTACAAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Freier SM;
                                                                                                                                                                                                                                                                              BP; 6 A;
                                                                                                                                                                                                                       Conservative
                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note= '
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             nucleotides"
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/*tag=
/mod_ba
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /*tag= a
/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note= "contains
linkages, all C 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           note= "2'-0-methoxyethyl nucleotide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       *tag=
                                                                                        DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         _base= OTHER
                                                                                                                                                                                                                                  1.9%;
                                                                                                                                                                                                                                                                              7 C; 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             87; 143pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "2'-O-methoxyethyl nucleotide"
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                                                                                        20
                                                                                                                                                                                                                      0;
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                                                                                                                                                                                                                                   Score 18.4;
Pred. No. 1.
                                                                                                                                                             20
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nucelotides are 5'-methylcytidine
                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                     1.3e+03
                                                                                                                                                                                                                                                                                0 Other;
                                                                                                                                                                                                                                                 DB 1; Length 20;
                                                                                                                                                                                                                       0
                                                                                                                                                                                                                      Gaps
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Human RANTES

oligonucleotide sequence

RESULT 774
ABZ89864/c
ID ABZ898
XX
AC ABZ898
XX
DT 17-OCT
XX
DE Human

ABZ89864 standard;

DNA;

20

17-OCT-2003 ABZ89864;

(first entry)

Human oligonucleotide sequence

밁 á

> 537 19;

CCTGCCTCAGCCTCCCAAGT 556

Conservative

<u>.</u>

Mismatches

Indels

0

Gaps

0,

ccreccreaeccreceaer 20

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CC immunosuppressive, and cytostatic active agent composition of the invention at a trianglammatory steroid and ubiquinone. A composition of the invention communosuppressive, antiallergic, antiasthmatic, hypotensive, an attinflammatory, antiallergic, antiasthmatic, hypotensive, are composition of the invention communosuppressive, and cytostatic activity. The composition may have a composition are respiratory, lung or malignant disease or condition, also composition are respiratory, lung or malignant disease or condition, also composition for reducing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of of or reducing sensitivity to adenosine, reducing levels of adenosine composition, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject stissue, or treating bronchoconstriction, increasing inflammation, lung allergies, or a respiratory disease or condition.

CC Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO are for whom the formation of the printed contribution is presented in the printed contribution.
Query Match
Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a novel pharmaceutical composition, which he first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking region; and 3' intron-exon junctions, or regions within 2-10 nucleotides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human, antisense, lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antia antiasthmatic; hypotensive; immunosuppressive; cytostatic; ge
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200285308-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Miller S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    24-APR-2001; 2001US-0286137P
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                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; SEQ ID NO 13207; 872pp;
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                                                                                                                                                                                                            ftp.wipo.int/pub/published_pct_sequences
                                       Similarity
                                                                                                                                               20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Li Y, Sa
Tang L,
                                                                                                                                            B₽;
                                                                                                                                            2 A;
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L, Shahabuddin
                                       1.9%;
                                                                                                                                            10 C; 4 G; 4 T; 0 U; 0 Other;
                                       Pred.
                                              Score 18.4;
Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    English.
                                              .3e+03;
                                                                                  DB 1;
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                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
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                                 17-OCT-2003
                                                                    ABZ97964;
                                                                                                      ABZ97964 standard;
                                                                                                                                                                                                                                                                                                                                         Sequence
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Tang
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                               (first entry)
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Shahabuddin
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Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
23-APR-2002; 2002WO-US013135.
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                                                                                                                                                                                                                                                     sapiens.
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corresponding RNAs, and Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or s Li Y, Sa Sandrasagra A, L, Shahabuddin Katz S;

24-APR-2001; 2001US-0286137P

(EPIG-) EPIGENESIS PHARM INC

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'n

Pabalan J,

Aguilar

SEQ ID NO 13206; 872pp; English

ubiquinone

nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO The invention relates to a novel pharmaceutical composition, which has first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or ftp.wipo.int/pub/published_pct_sequences

Sequence 20 BP; 2 A; 10 C; 2 G; 6 T; 0 U; 0 Other;

밁 S Matches Query Match Best Local S 532 19; r Similarity ATCCTCCTGCCTCAGCCTCC 551 ATTCTCCTGCCTCAGCCTCC Conservative 1.9%; 95.0%; 0 Score 18.4; Pred. No. 1. 20 Mismatches 1.3e+03; DB 1; Length 20; Indels 0 Gaps

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XXXXX RESULT 776 ABZ97909 17-OCT-2003 ABZ97909 standard; (first entry) DNA; 20

Human RANTES oligonucleotide sequence.

Human RANTES oligonucleotide sequence.

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RESULT 777
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
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                                                                                                            ABZ89861 standard;
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L, Shahabuddin S;
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                                                                                                            DNA;
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95.0%;
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Human oligonucleotide sequence

BXSXXX

17-OCT-2003 ABZ97902;

(first entry)

Human RANTES oligonucleotide sequence

ABZ97

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RESULT 778

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Matches Query Match Best Local (

Similarity

95.0%;

Conservative

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Sequence 20

BP; 4 A; 4 C; 9 G; 3 T; 0 U; 0 Other;

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Consistant active agent comprising an oligonucleotide antisense to the contribution relates to a novel pharmaceutical composition, which has a consistant coden, coding region, 5' or 3' end genomic flanking regions, coding region, 5' or 3' end genomic flanking regions, coding a polypeptide associated with lung and/or coding a polypeptide associated with lung and/or codinflammatory, steroid and thiquinone. A composition of the invention code antiinflammatory, antiallergic, antiasthmatic, hypotensive, and cycostatic activity. The composition may have a code in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also code for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing levels of adenosine code, or reducing sensitivity to adenosine, reducing levels of adenosine code, producing bronchodilation, increasing levels of building subject, lung surfactant in a subject, subject, or treating bronchoconstriction, lung surfactant in a subject, or a respiratory disease or condition. Composition in a subject, subject, or a respiratory disease or condition. The sequence data for this patent is not represented in the printed conspication, but was obtained in electronic format directly from WIPO and the printed of the printed of the printed or the printed or the printed of the printed of the printed of the printed or the 
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RESULT 779
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                                                                                                                  ABZ92724 standard; DNA;
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                                                                                                                                                                                                                                                  868
                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                  GGATTACAGGCGTGAGCCAC 887
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Tang
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                                                                                                                                                                                                                                                                                       Conservative
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L, Shahabuddin
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                                                                                                                                                                                                                                                                                                                                                              5 C; 8 G; 3 T; 0 U; 0 Other;
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                                                                                                                  BP.
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Pred. No. 1.
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Human oligonucleotide sequence

17-OCT-2003 ABZ92724;

(first entry)

RESULT 780 ABZ98012 ID ABZ980

standard;

DNA;

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4.75.

768 نر

TTTTTTGTATTTTTAGTAGA 787

TTTTTTGTATGTTTAGTAGA 20

Query Match Best Local S Matches 19

l Similarity 19; Conserv

Conservative

0;

1.98;

Score 18.4; DB 1; Length 20; Pred. No. 1.3e+03; O; Mismatches 1; Indels

0,

Gaps

0

Sequence

20 BP;

A; 0 C; 4

G; 12 T;

0 U;

0 Other;

XXXXXX

17-OCT-2003 ABZ98012; ABZ98012

(first entry)

Human RANTES oligonucleotide sequence.

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CC junctions of genes encoding a polypeptide associated with lung and/or CC nasal airway dysfunction and a second active agent comprising an CC antiinflammatory steroid and ubiquinone. A composition of the invention CC has antiinflammatory, antiallergic, antiasthmatic, hypotensive, and cytostatic activity. The composition may have a CC use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also CC for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of of or reducing sensitivity to adenosine, reducing levels of adenosine creceptor, producing bronchodilation, increasing levels of definition. CC lung surfactant in a subject's tissue, or treating bronchoconstriction. CC lung inflammation, lung allergies, or a respiratory disease or condition. CC lung inflammation, lung allergies, or a respiratory disease or condition. CC lung inflammation, but was obtained in electronic format directly from WIPO at figure in the printed constriction, but was obtained in electronic format directly from WIPO at figure is a figure of the sequence data for this patent is not represented in the printed constriction, but was obtained pot sequences.
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Miller S,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; SEQ ID NO 7966; 872pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 first active agent comprising an initiation codon, coding region, 5' and 3' intron-exon junctions,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          5' and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to a novel pharmaceutical
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CC first active agent comprising an oligonucleotide antisense to the CC initiation codon, coding region, 5 or 3 end genomic flanking regions, CC yand 3' intron-exon junctions, or regions within 2-10 nucleotides of CC junctions of genes encoding a polypeptide associated with lung and/or CC nasal airway dysfunction and a second active agent comprising an CC antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, CC immunosuppressive, and cytostatic activity. The composition may have a CC use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also CC for enhancing the prophylactic or therapeutic respiratory effect of an artiinflammatory steeroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine CC enterptor, producing bronchodilation, increasing levels of adenosine CC lung surfactant in a subject's tissue, or treating bronchoonstruction, CC lung inflammation, lung allergies, or a respiratory disease or condition. CC specification, but was obtained in electronic format directly from WIPO at ftp. wipo.int/pub/published_pct_sequences
RESULT 781
ABZ98015
ID ABZ980
XX
AC ABZ980
XX
DT 17-OCT
XX
DE Human
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
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                                                                                                                          ABZ98015;
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19; Conser
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, Tang L,
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L, Shahabuddin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1; Length 20;
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Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene thera
antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.
                                                                                                                                                                  gene therapy;
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Homo sapiens

WO200285308-A2

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P

(EPIG-) EPIGENESIS PHARM INC

Miller S Li Y, Tang ŗ Sandrasagra A, Shahabuddin S; Katz E, Pabalan Ģ Aguilar

U

WPI; 2003-229219/22

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone

Disclosure; SEQ ID NO 13257; 872pp; English

The invention relates to a novel pharmaceutical composition, which has a composition relates to a novel pharmaceutical composition, which has a composition codon, coding region, 5' or 3' end genomic flanking regions, coding region, 5' or 3' end genomic flanking regions, coding regions, coding regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or coding a polypeptide associated with lung and/or coding a polypeptide associated with lung and/or coding antiinflammatory steroid and ubiquinone. A composition of the invention code antiinflammatory, antiallergic, antiasthmatic, hypotensive, coding to a manusuppressive, and cytostatic activity. The composition may have a composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also compositing a respiratory, lung or malignant disease or condition, also composition generating the prophylactic or therapeutic respiratory effect of an antialnflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine contractant in a subject, for activity reducing levels of subiquinone or receptor, producing bronchodilation, increasing levels of subiquinone or anniant contractant in a subject of the composition of the composition and contractant in a subject, for reducing levels of subiquinone or anniant contractant co lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO ftp.wipo.int/pub/published_pct_sequences

Sequence 20 BP; 3 A; 4 C; 9 G; 4 T; 0 U; 0 Other;

Matches Query Match Local 19; Similarity Conservative 1.9%; 0, Score 18.4; Pred. No. 1. Mismatches .3e+03; DB 1; Length 20; Indels 0 Gaps

0

뭉 Ś 651 GGAGTGCAGTGGCGCAATCT 670 Н GGAGTGCAGTGGCGCGATCT 20

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RESULT 782 ABZ97908 standard; DNA;

ΒP

XXXXXXX 17-OCT-2003 (first entry)

ABZ97908

Human RANTES oligonucleotide sequence

Human RANTES

oligonucleotide (first entry)

sequence

17-OCT-2003

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RESULT 783
ABZ98003
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Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at fire wino int/puh/muhlished for sequences.
                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 19; Conserv
Human RANTES oligonucleotide sequence.
                                         17-OCT-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
                                                                              ABZ98003
                                                                                                                   ABZ98003 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                Sequence 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (EPIG-) EPIGENESIS PHARM INC
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                                                                                                                                                                                                                                                    CTCAGCCTCCCAAGTAGCTG 561
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Tang L,
                                                                                                                                                                                                                                                                                                                                                                BP; 3 A; 8 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                         Conservative
                                         (first entry)
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L, Shahabuddin
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                                                                                                                                                                                                                                                                                                         1.9%;
                                                                                                                   ВP
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Pred. No. 1
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                                                                                                                                                                                                                                                                                         Mismatches
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RESULT 784
ABZ97903
ID ABZ979
XX
AC ABZ979
XX
DT 17-OCT
XX
DE Human

ABZ97903;

ABZ97903 standard; DNA;

20 BP

17-OCT-2003

(first entry)

Human RANTES oligonucleotide sequence.

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20

Matches

19;

642

ACCCAGGCTGGAGTGCAGTG 661 Conservative

Query Match Best Local Similarity

95.0%;

Score 18.4; Pred. No. 1.

.3e+03 DB 1;

Length

<u>,,</u>

Mismatches

0

Gaps

0

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initiation codon, coding region, 5' or 3' end genomic flanking regions, CC 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of cyunctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory, antiallergic, antiasthmatic, hypotensive, and cytostatic activity. The composition may have a cure in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also conformation the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or adenosine receptor, producing bronchodilation, increasing levels of adenosine reducing inflammation, lung allergies, or a respiratory disease or condition. CC lung inflammation, lung allergies, or a respiratory disease or condition. Solve: The sequence data for this patent is not represented in the printed generation, but was obtained in electronic format directly from WIPO or a fer wino int/nub/wiblished not segmences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
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Sequence 20 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; SEQ ID NO 13245; 872pp; English
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                                                                              ftp.wipo.int/pub/published_pct_sequences
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5 A; 4 C; 8
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L, Shahabuddin
G; 3 T; 0 U; 0 Other;
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RESULT 785
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ID ABZ990
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DT 17-OCT
XX
DE Human
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
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    Human PDE4C oligonucleotide sequence
                                          17-OCT-2003
                                                                                                                    ABZ99062 standard;
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19; Conser
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, Tang L,
                                                                                                                                                                                                                                                       ACAGGCGTGAGCCACCACGC 892
                                          (first entry)
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L, Shahabuddin
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Human, antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy, antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.
                                                                                                                                                                                therapy;
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Homo sapiens
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WO200285308-A2

23-APR-2002; 2002WO-US013135

24-APR-2001; 2001US-0286137P

(EPIG-) EPIGENESIS PHARM INC

Miller ŝ Li Y, Sa Tang L, Sandrasagra A, Shahabuddin S Katz E, Pabalan ŗ Aguilar

WPI; 2003-229219/22

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone

Disclosure; SEQ ID NO 14304; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a cfirst active agent comprising an oligonucleotide antisense to the cinitiation codon, coding region, 5' or 3' end genomic flanking regions, 5' or 3' end genomic flanking regions, 6' or and 3' intron-exon junctions, or regions within 2-10 nucleotides of cymctions of genes encoding a polypeptide associated with lung and/or comparison and a second active agent comprising an capture antiinflammatory steroid and ubiquinone. A composition of the invention capture in antisense gene therapy. The composition may have a cuse in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also capture in a subject, for reducing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing levels of adenosine capturing a respiratory to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of bulguinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, continuing antiflammation, lung allargies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at the manner of the printed capture in the printed specification, but was obtained in electronic format directly from WIPO at the manner of the printed capture in the printed specification. ftp.wipo.int/pub/published_pct_sequences

Sequence 20 BP; 3 A; 10 C; 3 G; 4 T; 0 U; 0 Other;

밁 á Matches Query Match Best Local 538 19; Similarity CTGCCTCAGCCTCCCAAGTA 557 CTGCCTCAGCCTCCCCAGTA 20 Conservative 1.9%; 0 Score 18.4; Pred. No. 1. Mismatches .3e+03; DB 1; Length Indels 20; 0, Gaps

0

RESULT 786 ABZ99105; ABZ99105 standard; DNA; 20

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HXHXAX 17-OCT-2003 (first entry)

Human PDE4C oligonucleotide sequence.

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BXAXAXB
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ABZ98013
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Best Local Similarity
Matches 19; Conserv
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Miller S
                                    17-OCT-2003
                                                                           ABZ98013;
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                                                                                                                ABZ98013 standard; DNA;
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L, Shahabuddin
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Human RANTES oligonucleotide sequence.

EXE

17-OCT-2003

(first entry)

Human PDE4C oligonucleotide sequence.

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RESULT 788 ABZ99071

ABZ99071

standard; DNA;

片 S

μ

CGCCCAGGCTGGAGTGCAGT 20

641 CACCCAGGCTGGAGTGCAGT 660

Matches Query Match Best Local :

19;

Conservative

0

Mismatches

Indels

<u>,,</u>

Gaps

0

Similarity

1.9**%**; 95.0**%**;

Score 18.4; Pred. No. 1.

.3e+03

DB 1; Length 20;

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cc mass airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention chas antiinflammatory, antiallergic, antiasthmatic, hypotensive, cc immunosuppressive, and cytostatic activity. The composition may have a composition may have a composition are spiratory, lung or malignant disease or condition, also cc for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of of or reducing sensitivity to adenosine, reducing levels of adenosine ceeptor, producing bronchodilation, increasing levels of adenosine colung surfactant in a subject's tissue, or treating bronchoconstriction. Cc lung inflammation, lung allergies, or a respiratory disease or condition. Cc lung inflammation, lung allergies, or a respiratory disease or condition. Cc lung inflammation, but was obtained in electronic format directly from WIPO cc at ftp.wipo.int/pub/published_pct_sequences
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Miller S
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antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic;
antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy
antisense gene therapy; respiratory; lung; adenosine sensitivity;
adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a novel pharmaceutical composition, which has first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5 or 3' end genomic flanking region, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or
   Sequence
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Tang
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L, Shahabuddin
A; 6 C; 8
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   G; 3 T; 0
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   U; 0 Other;
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RESULT 789
ABZ89844/c
ID ABZ898
XX
AC ABZ898
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DT 17-OCT
XX
DE Human 0
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Tang
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                                                                                      (first entry)
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L, Shahabuddin
                                                                                                                                                                                                                                            DNA;
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Pred. No. 1.
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antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.
                                                                                                                     Human, antisense, lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antia antiasthmatic; hypotensive; immunosuppressive; cytostatic; ge
                                                                                                                                                                                antiallergic;
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Homo sapiens.
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WO200285308-A2

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P

(EPIG-) EPIGENESIS PHARM INC

Miller ģ Li Y, Sa Tang L, Sandrasagra A, L, Shahabuddin ŝ Katz E, Pabalan J, Aguilar

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or

Disclosure; SEQ ID NO 5086; 872pp; English

cy junctions of gene encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention cantiinflammatory, antiallergic, antiasthmatic, hypotensive, has antiinflammatory, antiallergic, antiasthmatic, hypotensive, and cytostatic activity. The composition may have a constitution of the invention of the invention of the invention of the prophylactic or therapy. The composition may have a preventing a respiratory, lung or malignant disease or condition, also containflammatory steroid in a subject, for reducing or depleting levels of an antiinflammatory steroid in a subject, for reducing or depleting levels of confort, producing bronchodilation, increasing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine confort inflammation, lung allergies, or respiratory disease or condition. CC lung surfactant in a subject's tissue, or treating bronchoconstriction, content of the sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO appears to the printed of the printe The invention relates to a novel pharmaceutical composition, which has first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, initiation codon, coding region, 5' or 3' end genomic flanking regions 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of ftp.wipo.int/pub/published_pct_sequences

Sequence 20 BP; 13 A; 2 C; 0 G; 5 T; 0 U; 0 Other;

밁 S ABZ92736 Matches Query Match ABZ92736 standard; DNA; Local 20 19; Similarity ATTTTTTTGTATTTTTAGTA 785 Conservative 1.9%; 먪 0; Score 18.4; Pred. No. 1 Mismatches .3e+03 DB 1; Length Indels 0 Gaps

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Human oligonucleotide sequence

BXAXAXU Human oligonucleotide sequence 17-OCT-2003 ABZ92736; (first entry)

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RESULT 791
ABZ99077
ID ABZ990
XX
AC ABZ990
XX
DT 17-OCT
XX
DE Human
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Best Local S
Matches 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20
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Tang
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L, Shahabuddin
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Pred. No. 1.3e+03;
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밁 S

Human PDE4C oligonucleotide sequence.

17-OCT-2003

(first entry)

RESULT 792
ACA88928/c
ID ACA889
XX
AC ACA889
XX
OT 08-JUL
XX
DE Select

08-JUL-2003 ACA88928;

(first entry)

Selection and amplification of genetic markers PCR related primer #39

ACA88928 standard; DNA;

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Matches Query Match

389 19;

AAAGTGCTGGGATTACAGGC 408 AAAGTGCTGGGATTATAGGC 20

Local

Similarity

Conservative

<u>.</u>

Score 18.4; Pred. No. 1. Mismatches

4; DB 1; 1.3e+03; 0 Other;

Length 20 Indels

0,

Gaps

0

Sequence

20 BP; 6

A; 2 C; 7 1.9%;

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CC first active agent comprising an oligonucleotide antisense to the cinitiation codon, coding region, 5' or 3' end genomic flanking regions, CC 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of CC junctions of genes encoding a polypeptide associated with lung and/or CC nasal airway dysfunction and a second active agent comprising an CC antiinflammatory steroid and ubiquinone. A composition of the invention CC has antiinflammatory, antiallergic, antiasthmatic, hypotensive, CC immunosuppressive, and cytostatic activity. The composition may have a CC use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also CC for enhancing the prophylactic or therapeutic respiratory effect of an CC antiinflammatory steroid in a subject, for reducing or depleting levels of of, or reducing sensitivity to adenosine, reducing levels of adenosine caceptor, producing bronchodilation, increasing levels of ubiquinone or CC lung inflammation, lung allergise, or a respiratory disease or condition. CC Note: The sequence data for this patent is not represented in the printed of a precification, but was obtained in electronic format directly from WIPO at frem wino information accounts.
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Miller S,
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antisense gene therapy; respiratory; lung; adenosine sensitivity;
adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
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at ftp.wipo.int/pub/published_pct_sequences
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